

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
8 March 2001 (08.03.2001)

PCT

(10) International Publication Number
WO 01/16312 A2

(51) International Patent Classification⁷: **C12N 15/11**,
9/00, C07H 21/00, 19/00, C12P 19/34, A61K 31/7088,
C12N 5/10 // A61P 3/10, 9/00, 25/28, 35/00

(21) International Application Number: PCT/US00/23998

(22) International Filing Date: 30 August 2000 (30.08.2000)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:

60/151,713	31 August 1999 (31.08.1999)	US
09/406,643	27 September 1999 (27.09.1999)	US
60/156,467	27 September 1999 (27.09.1999)	US
60/156,236	27 September 1999 (27.09.1999)	US
09/436,430	8 November 1999 (08.11.1999)	US
60/169,100	6 December 1999 (06.12.1999)	US
60/173,612	29 December 1999 (29.12.1999)	US
09/474,432	29 December 1999 (29.12.1999)	US
09/476,387	30 December 1999 (30.12.1999)	US
09/498,824	4 February 2000 (04.02.2000)	US
09/531,025	20 March 2000 (20.03.2000)	US
60/197,769	14 April 2000 (14.04.2000)	US
09/578,223	23 May 2000 (23.05.2000)	US
Not furnished	9 August 2000 (09.08.2000)	US

(71) Applicant (for all designated States except US): **RI-BOZYME PHARMACEUTICALS, INC.** [US/US];
2950 Wilderness Place, Boulder, CO 80301 (US).

(72) Inventors; and

(75) Inventors/Applicants (for US only): **MCSWIGGEN, James** [US/US]; 4866 Franklin Drive, Boulder, CO 80301 (US). **USMAN, Nassim** [CA/US]; 2129 Night Sky Lane, Lafayette, CO 80026 (US). **BLATT, Lawrence** [US/US]; 2176 Riverside Lane, Boulder, CO 80304 (US). **BEIGEL-MAN, Leonid** [US/US]; 5530 Colt Drive, Longmont, CO 80503 (US). **BURGIN, Alex** [US/US]; 832 Caminito Estrella, Chula Vista, CA 91910 (US). **KARPEISKY,**

Alexander [RU/US]; 420 Vernier Avenue, Lafayette, CO 80026 (US). **MATULIC-ADAMIC, Jasenka** [HR/US]; 760 South 42nd Street, Boulder, CO 80303 (US). **SWEEDLER, David** [US/US]; 956 St. Andrews Lane, Louisville, CO 80027 (US). **DRAPER, Kenneth** [US/US]; 4791 Cougar Creek Trail, Reno, NV 89509 (US). **CHOWRIRA, Bharat** [IN/US]; 1138 Clubhouse Drive, Broomfield, CO 80020 (US). **STINCHCOMB, Dan** [US/US]; 8409 South Country Road 3, Fort Collins, CO 80528 (US). **BEAUDRY, Amber** [US/US]; 13068 Westlake Place, Broomfield, CO 80026 (US). **ZINNEN, Shawn** [US/US]; 2378 Birch Street, Denver, CO 80207 (US). **LUGWIG, Janos** [DE/DE]; Untere Karspule 13B, D-37073 Gottingen (DE). **SPROAT, Brian, S.** [GB/DE]; Am Antonsberg 10, D-37139 Adelebsen (DE).

(74) Agents: **PRINCE, Robert, W.** et al.; Brobeck, Phleger & Harrison, 12390 El Camino Real, San Diego, CA 92130 (US).

(81) Designated States (national): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW.

(84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).

Published:

— Without international search report and to be republished upon receipt of that report.

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

WO 01/16312 A2

(54) Title: NUCLEIC ACID BASED MODULATORS OF GENE EXPRESSION

(57) Abstract: Novel nucleic acid molecules useful as inhibitors of gene expression, compositions, and methods for their use.

NUCLEIC ACID BASED MODULATORS OF GENE EXPRESSION

Background of the Invention

This invention relates to reagents useful as inhibitors of gene expression relating to diseases such as cancers, diabetes, obesity, Alzheimer's disease, cardiac diseases, age-related diseases, and/or hepatitis B infections and related conditions.

Summary of the Invention

The invention features novel nucleic acid-based techniques [e.g., enzymatic nucleic acid molecules (ribozymes), antisense nucleic acids, 2-5A antisense chimeras, triplex DNA, antisense nucleic acids containing RNA cleaving chemical groups (for example, Cook et al., U.S. Patent 5,359,051)] and methods for their use to modulate the expression of molecular targets impacting the development and progression of cancers, diabetes, obesity, Alzheimer's disease, cardiac diseases, age-related diseases, and/or hepatitis B infections and related conditions

In a preferred embodiment, the invention features novel nucleic acid-based techniques [e.g., enzymatic nucleic acid molecules (ribozymes), antisense nucleic acids, 2-5A antisense chimeras, triplex DNA, antisense nucleic acids containing RNA cleaving chemical groups (for example, Cook et al., U.S. Patent 5,359,051)] and methods for their use for inhibiting the expression of disease related genes, e.g., Protein-Tyrosine-Phosphatase-1b (PTP-1B, Genbank accession No. NM_002827), Methionine Aminopeptidase (MetAP-2, Genbank accession No. U29607), beta-Secretase (BACE, Genbank accession No. AF190725), Presenilin-1 (ps-1, Genbank accession No. L76517), Presenilin-2 (ps-2, Genbank accession No. L43964), Human Epidermal Growth Factor Receptor-2 (HER2/c-erb2/neu, Genbank accession No. X03363), Phospholamban (PLN, Genbank accession No. NM_002667), Telomerase (TERT, Genbank accession No. NM_003219) and Hepatitis B virus genes (HBV, Genbank accession No. AF100308.1). Such ribozymes can be used in a method for treatment of diseases caused by the expression of these genes in man and other animals, including other primates.

Thus, in an additional preferred embodiment, the invention features novel nucleic acid-based techniques such as enzymatic nucleic acid molecules and antisense molecules and methods for their use to down regulate or inhibit the expression of genes encoding Protein-Tyrosine-Phosphatase-1b (PTP-1B), Methionine Aminopeptidase (MetAP-2),

beta-Secretase (BACE), Presenilin-1 (ps-1), Presenilin-2 (ps-2), Human Epidermal Growth Factor Receptor-2 (HER2/c-erb2/neu), Phospholamban (PLN), Telomerase (hTERT) PKC alpha. and Hepatitis B (HBV) proteins. In particular, applicant describes the selection and function of nucleic acid molecules capable of cleaving RNAs encoded by these genes and their use to reduce levels of PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV proteins in various tissues to treat the diseases discussed herein. Such nucleic acid molecules are also useful for diagnostic uses.

In a preferred embodiment, the invention features the use of one or more of the nucleic acid-based techniques independently or in combination to inhibit the expression of the genes encoding PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV. Specifically, the invention features the use of nucleic acid-based techniques to specifically inhibit the expression of PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, PKC alpha, and/or HBV genes.

In yet another preferred embodiment, the invention features the use of an enzymatic nucleic acid molecule, preferably in the hammerhead, NCH (Inozyme), G-cleaver, amberzyme, zinzyme, and/or DNAzyme motif, to inhibit the expression of PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, PKC alpha and/or HBV RNA.

Applicant indicates that these nucleic acid molecules are able to inhibit expression of PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, PKC alpha, and/or HBV genes. Those of ordinary skill in the art, will find that it is clear from the examples described that other nucleic acid molecules that inhibit target PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV encoding mRNAs may be readily designed and are within the scope of the invention.

By "inhibit" it is meant that the activity of target genes or level of mRNAs or equivalent RNAs encoding target genes is reduced below that observed in the absence of the nucleic acid molecules of the instant invention (*e.g.*, enzymatic nucleic acid molecules), antisense nucleic acids, 2-5A antisense chimeras, triplex DNA, antisense nucleic acids containing RNA cleaving chemical groups). In one embodiment, inhibition with an enzymatic nucleic acid molecule preferably is below that level observed in the presence of an enzymatically attenuated nucleic acid molecule that is able to bind to the same site on the mRNA, but is unable to cleave that RNA. In another embodiment, inhibition with nucleic acid molecules, including enzymatic nucleic acid and antisense

molecules, is preferably greater than that observed in the presence of, for example, an oligonucleotide with scrambled sequence or with mismatches. In another embodiment, inhibition of target genes with the nucleic acid molecule of the instant invention is greater than in the presence of the nucleic acid molecule than in its absence. According to the invention, the activity of telomerase enzyme or the level of RNA encoding one or more protein subunits of the telomerase enzyme is inhibited if it is at least 10% less, 20% less, 50% less, 75% less or even not active or present at all, in the presence of a nucleic acid of the invention relative to the level in the absence of such a nucleic acid.

By "enzymatic nucleic acid molecule" it is meant a nucleic acid molecule which has complementarity in a substrate binding region to a specified gene target, and also has an enzymatic activity which is active to specifically cleave target RNA. That is, the enzymatic nucleic acid molecule is able to intermolecularly cleave RNA and thereby inactivate a target RNA molecule. These complementary regions allow sufficient hybridization of the enzymatic nucleic acid molecule to the target RNA and thus permit cleavage. One hundred percent complementarity is preferred, but complementarity as low as 50-75% may also be useful in this invention. The nucleic acids may be modified at the base, sugar, and/or phosphate groups. The term enzymatic nucleic acid is used interchangeably with phrases such as ribozymes, catalytic RNA, enzymatic RNA, catalytic DNA, aptazyme or aptamer-binding ribozyme, regulatable ribozyme, catalytic oligonucleotides, nucleozyme, DNAzyme, RNA enzyme, endoribonuclease, endonuclease, minizyme, leadzyme, oligozyme or DNA enzyme. All of these terminologies describe nucleic acid molecules with enzymatic activity. The specific enzymatic nucleic acid molecules described in the instant application are not meant to be limiting and those skilled in the art will recognize that all that is important in an enzymatic nucleic acid molecule of this invention is that it have a specific substrate binding site which is complementary to one or more of the target nucleic acid regions, and that it have nucleotide sequences within or surrounding that substrate binding site which impart a nucleic acid cleaving activity to the molecule (Cech et al., U.S. Patent No. 4,987,071; Cech et al., 1988, JAMA 260:20 3030-4).

By "nucleic acid molecule" as used herein is meant a molecule having nucleotides. The nucleic acid can be single, double, or multiple stranded and may comprise modified or unmodified nucleotides or non-nucleotides or various mixtures and combinations thereof.

An example of a nucleic acid molecule according to the invention is a gene which encodes for a macromolecule such as a protein.

By "enzymatic portion" or "catalytic domain" is meant that portion/region of the enzymatic nucleic acid molecule essential for cleavage of a nucleic acid substrate (for example see **Figures 1-5**).

By "substrate binding arm" or "substrate binding domain" is meant that portion/region of a ribozyme which is complementary to (*i.e.*, able to base-pair with) a portion of its substrate. Generally, such complementarity is 100%, but can be less if desired. For example, as few as 10 bases out of 14 may be base-paired. Such arms are shown generally in **Figures 1-5**. That is, these arms contain sequences within a ribozyme which are intended to bring ribozyme and target RNA together through complementary base-pairing interactions. The ribozyme of the invention may have binding arms that are contiguous or non-contiguous and may be of varying lengths. The length of the binding arm(s) are preferably greater than or equal to four nucleotides and of sufficient length to stably interact with the target RNA; specifically 12-100 nucleotides; more specifically 14-24 nucleotides long. If two binding arms are chosen, the design is such that the length of the binding arms are symmetrical (*i.e.*, each of the binding arms is of the same length; *e.g.*, five and five nucleotides, six and six nucleotides or seven and seven nucleotides long) or asymmetrical (*i.e.*, the binding arms are of different length; *e.g.*, six and three nucleotides; three and six nucleotides long; four and five nucleotides long; four and six nucleotides long; four and seven nucleotides long; and the like). Binding arms can be complementary to the specified substrate, to a portion of the indicated substrate, to the indicated substrate sequence and additional adjacent sequence, or a portion of the indicated sequence and additional adjacent sequence.

By "NCH" or "Inozyme" motif is meant, an enzymatic nucleic acid molecule comprising a motif as described in Ludwig *et al.*, USSN No. 09/406,643, filed September 27, 1999, entitled "COMPOSITIONS HAVING RNA CLEAVING ACTIVITY", and International PCT publication Nos. WO 98/58058 and WO 98/58057, all incorporated by reference herein in their entirety, including the drawings.

By "G-cleaver" motif is meant, an enzymatic nucleic acid molecule comprising a motif as described in Eckstein *et al.*, International PCT publication No. WO 99/16871, incorporated by reference herein in its entirety, including the drawings.

By "zinzyme" motif is meant, a class II enzymatic nucleic acid molecule comprising a motif as described herein and in Beigelman *et al.*, International PCT publication No. WO 99/55857, incorporated by reference herein in its entirety, including the drawings.

5 By "amberzyme" motif is meant, a class I enzymatic nucleic acid molecule comprising a motif as described herein and in Beigelman *et al.*, International PCT publication No. WO 99/55857, incorporated by reference herein in its entirety, including the drawings.

By 'DNAzyme' is meant, an enzymatic nucleic acid molecule lacking a ribonucleotide (2'-OH) group. In particular embodiments, the enzymatic nucleic acid molecule may have an attached linker(s) or other attached or associated groups, moieties, or chains containing one or more nucleotides with 2'-OH groups. A DNAzyme can be synthesized chemically or can be expressed by means of a single stranded DNA vector or equivalent thereof.

15 By "sufficient length" is meant an oligonucleotide of greater than or equal to 3 nucleotides that is of a length great enough to provide the intended function under the expected condition. For example, for binding arms of enzymatic nucleic acid "sufficient length" means that the binding arm sequence is long enough to provide stable binding to a target site under the expected binding conditions. Preferably, the binding arms are not so long as to prevent useful turnover.

20 By "stably interact" is meant, interaction of the oligonucleotides with target nucleic acid (*e.g.*, by forming hydrogen bonds with complementary nucleotides in the target under physiological conditions).

By "equivalent" RNA to PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV is meant to include those naturally occurring RNA molecules having homology (partial or complete) to PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV proteins or encoding for proteins with similar function as PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV in various organisms, including human, rodent, primate, rabbit, pig, protozoans, fungi, plants, and other microorganisms and parasites. The equivalent RNA sequence also includes in addition to the coding region, regions such as 5'-untranslated region, 3'-untranslated region, introns, intron-exon junction and the like in HBV.

By "homology" is meant the nucleotide sequence of two or more nucleic acid molecules is partially or completely identical.

By "antisense nucleic acid", it is meant a non-enzymatic nucleic acid molecule that binds to target RNA by means of RNA-RNA or RNA-DNA or RNA-PNA (protein nucleic acid; Egholm *et al.*, 1993 *Nature* 365, 566) interactions and alters the activity of the target RNA (for a review, see Stein and Cheng, 1993 *Science* 261, 1004 and Woolf *et al.*, US patent No. 5,849,902). Typically, antisense molecules will be complementary to a target sequence along a single contiguous sequence of the antisense molecule. However, in certain embodiments, an antisense molecule may bind to substrate such that the substrate molecule forms a loop, and/or an antisense molecule may bind such that the antisense molecule forms a loop. Thus, the antisense molecule may be complementary to two (or even more) non-contiguous substrate sequences or two (or even more) non-contiguous sequence portions of an antisense molecule may be complementary to a target sequence or both. For a review of current antisense strategies, see Schmajuk *et al.*, 1999, *J. Biol. Chem.*, 274, 21783-21789, Delihias *et al.*, 1997, *Nature*, 15, 751-753, Stein *et al.*, 1997, *Antisense N. A. Drug Dev.*, 7, 151, Crooke, 1998, *Biotech. Genet. Eng. Rev.*, 15, 121-157, Crooke, 1997, *Ad. Pharmacol.*, 40, 1-49. In addition, antisense DNA can be used to target RNA by means of DNA-RNA interactions, thereby activating RNase H, which digests the target RNA in the duplex. Antisense DNA can be synthesized chemically or can be expressed via the use of a single stranded DNA expression vector or the equivalent thereof.

By "2-5A antisense chimera" it is meant, an antisense oligonucleotide containing a 5'-phosphorylated 2'-5'-linked adenylyate residue. These chimeras bind to target RNA in a sequence-specific manner and activate a cellular 2-5A-dependent ribonuclease which, in turn, cleaves the target RNA (Torrence *et al.*, 1993 *Proc. Natl. Acad. Sci. USA* 90, 1300).

By "triplex DNA" it is meant an oligonucleotide that can bind to a double-stranded DNA in a sequence-specific manner to form a triple-strand helix. Formation of such triple helix structure has been shown to inhibit transcription of the targeted gene (Duval-Valentin *et al.*, 1992, *Proc. Natl. Acad. Sci. USA*, 89, 504).

By "gene" it is meant a nucleic acid that encodes a RNA.

By "complementarity" is meant that a nucleic acid can form hydrogen bond(s) with another RNA sequence by either traditional Watson-Crick or other non-traditional types. In reference to the nucleic molecules of the present invention, the binding free energy for a nucleic acid molecule with its target or complementary sequence is sufficient to allow the relevant function of the nucleic acid to proceed, e.g., ribozyme cleavage, antisense or triple helix inhibition. Determination of binding free energies for nucleic acid molecules is well known in the art (see, e.g., Turner et al., 1987, *CSH Symp. Quant. Biol.* LII pp.123-133; Frier et al., 1986, *Proc. Nat. Acad. Sci. USA* 83:9373-9377; Turner et al., 1987, *J. Am. Chem. Soc.* 109:3783-3785). A percent complementarity indicates the percentage of contiguous residues in a nucleic acid molecule which can form hydrogen bonds (e.g., Watson-Crick base pairing) with a second nucleic acid sequence (e.g., 5, 6, 7, 8, 9, 10 out of 10 being 50%, 60%, 70%, 80%, 90%, and 100% complementary). "Perfectly complementary" means that all the contiguous residues of a nucleic acid sequence will hydrogen bond with the same number of contiguous residues in a second nucleic acid sequence.

At least seven basic varieties of naturally-occurring enzymatic RNAs are known presently. Each can catalyze the hydrolysis of RNA phosphodiester bonds in *trans* (and thus can cleave other RNA molecules) under physiological conditions. **Table I** summarizes some of the characteristics of these ribozymes. In general, enzymatic nucleic acids act by first binding to a target RNA. Such binding occurs through the target binding portion of a enzymatic nucleic acid which is held in close proximity to an enzymatic portion of the molecule that acts to cleave the target RNA. Thus, the enzymatic nucleic acid first recognizes and then binds a target RNA through complementary base-pairing, and once bound to the correct site, acts enzymatically to cut the target RNA. Strategic cleavage of such a target RNA will destroy its ability to direct synthesis of an encoded protein. After an enzymatic nucleic acid has bound and cleaved its RNA target, it is released from that RNA to search for another target and can repeatedly bind and cleave new targets. Thus, a single ribozyme molecule is able to cleave many molecules of target RNA. In addition, the ribozyme is a highly specific inhibitor of gene expression, with the specificity of inhibition depending not only on the base-pairing mechanism of binding to the target RNA, but also on the mechanism of target RNA cleavage. Single mismatches,

or base-substitutions, near the site of cleavage can completely eliminate catalytic activity of a ribozyme.

The enzymatic nucleic acid molecule that cleave the specified sites in PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV-specific RNAs represent a novel therapeutic approach to treat a variety of pathologic indications, including, HBV infection, hepatitis, hepatocellular carcinoma, tumorigenesis, cirrhosis, liver failure, cancers including breast, ovarian, prostate, and esophageal cancer, tumorigenesis, retinopathy, arthritis, psoriasis, female reproduction, restinosis, certain infectious diseases, transplant rejection and autoimmune disease such as multiple sclerosis, lupus, and AIDS, age related diseases such as macular degeneration and skin ulceration, Alzheimer's disease, dementia, diabetes, obesity and any other condition related to the level of PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV in a cell or tissue.

In one of the preferred embodiments of the inventions described herein, the enzymatic nucleic acid molecule is formed in a hammerhead or hairpin motif, but may also be formed in the motif of a hepatitis delta virus, group I intron, group II intron or RNase P RNA (in association with an RNA guide sequence), *Neurospora* VS RNA, DNAzymes, NCH cleaving motifs, or G-cleavers. Examples of such hammerhead motifs are described by Dreyfus, *supra*, Rossi *et al.*, 1992, *AIDS Research and Human Retroviruses* 8, 183. Examples of hairpin motifs are described by Hampel *et al.*, EP0360257, Hampel and Tritz, 1989 *Biochemistry* 28, 4929, Feldstein *et al.*, 1989, *Gene* 82, 53, Haseloff and Gerlach, 1989, *Gene*, 82, 43, Hampel *et al.*, 1990 *Nucleic Acids Res.* 18, 299; and Chowrira & McSwiggen, US. Patent No. 5,631,359. The hepatitis delta virus motif is described by Perrotta and Been, 1992 *Biochemistry* 31, 16. The RNase P motif is described by Guerrier-Takada *et al.*, 1983 *Cell* 35, 849; Forster and Altman, 1990, *Science* 249, 783; and Li and Altman, 1996, *Nucleic Acids Res.* 24, 835. The *Neurospora* VS RNA ribozyme motif is described by Collins (Saville and Collins, 1990 *Cell* 61, 685-696; Saville and Collins, 1991 *Proc. Natl. Acad. Sci. USA* 88, 8826-8830; Collins and Olive, 1993 *Biochemistry* 32, 2795-2799; and Guo and Collins, 1995, *EMBO. J.* 14, 363). Group II introns are described by Griffin *et al.*, 1995, *Chem. Biol.* 2, 761; Michels and Pyle, 1995, *Biochemistry* 34, 2965; and Pyle *et al.*, International PCT Publication No. WO 96/22689. The Group I intron is described by Cech *et al.*, U.S. Patent 4,987,071. DNAzymes are described by Usman *et al.*, International PCT Publication No. WO 95/11304; Chartrand *et*

al., 1995, *NAR* 23, 4092; Breaker *et al.*, 1995, *Chem. Bio.* 2, 655; and Santoro *et al.*, 1997, *PNAS* 94, 4262. NCH cleaving motifs are described in Ludwig & Sproat, International PCT Publication No. WO 98/58058; and G-cleavers are described in Kore *et al.*, 1998, *Nucleic Acids Research* 26, 4116-4120 and Eckstein *et al.*, International PCT Publication No. WO 99/16871. Additional motifs include the Aptazyme (Breaker *et al.*, WO 98/43993), Amberzyme (Class I motif; Figure 3; Beigelman *et al.*, International PCT publication No. WO 99/55857) and Zinzyme (Beigelman *et al.*, International PCT publication No. WO 99/55857), all these references are incorporated by reference herein in their totalities, including drawings and can also be used in the present invention. These specific motifs are not limiting in the invention and those skilled in the art will recognize that all that is important in an enzymatic nucleic acid molecule of this invention is that it has a specific substrate binding site which is complementary to one or more of the target gene RNA regions, and that it have nucleotide sequences within or surrounding that substrate binding site which impart an RNA cleaving activity to the molecule (Cech *et al.*, U.S. Patent No. 4,987,071).

In preferred embodiments of the present invention, a nucleic acid molecule, *e.g.*, an antisense molecule, a triplex DNA, or a ribozyme, is 13 to 100 nucleotides in length, *e.g.*, in specific embodiments 35, 36, 37, or 38 nucleotides in length (*e.g.*, for particular ribozymes or antisense). In particular embodiments, the nucleic acid molecule is 15-100, 17-100, 20-100, 21-100, 23-100, 25-100, 27-100, 30-100, 32-100, 35-100, 40-100, 50-100, 60-100, 70-100, or 80-100 nucleotides in length. Instead of 100 nucleotides being the upper limit on the length ranges specified above, the upper limit of the length range can be, for example, 30, 40, 50, 60, 70, or 80 nucleotides. Thus, for any of the length ranges, the length range for particular embodiments has lower limit as specified, with an upper limit as specified which is greater than the lower limit. For example, in a particular embodiment, the length range can be 35-50 nucleotides in length. All such ranges are expressly included. Also in particular embodiments, a nucleic acid molecule can have a length which is any of the lengths specified above, for example, 21 nucleotides in length.

In a preferred embodiment, the invention provides a method for producing a class of nucleic acid-based gene inhibiting agents which exhibit a high degree of specificity for the RNA of a desired target. For example, the enzymatic nucleic acid molecule is preferably targeted to a highly conserved sequence region of target RNAs encoding PTP-1B, MetAP-

2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV proteins (specifically PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV RNA) such that specific treatment of a disease or condition can be provided with either one or several nucleic acid molecules of the invention. Such nucleic acid molecules can be delivered exogenously to specific tissue or cellular targets as required. Alternatively, the nucleic acid molecules (e.g., ribozymes and antisense) can be expressed from DNA and/or RNA vectors that are delivered to specific cells.

As used in herein "cell" is used in its usual biological sense, and does not refer to an entire multicellular organism, e.g., specifically does not refer to a human. The cell may be present in an organism which may be a human but is preferably a non-human multicellular organism, e.g., birds, plants and mammals such as cows, sheep, apes, monkeys, swine, dogs, and cats. The cell may be prokaryotic (e.g., bacterial cell) or eukaryotic (e.g., mammalian or plant cell).

By "PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV proteins" is meant, a protein or a mutant protein derivative thereof, comprising sequence expressed and/or encoded by PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, genes and/or the HBV genome respectively.

By "highly conserved sequence region" is meant a nucleotide sequence of one or more regions in a target gene does not vary significantly from one generation to the other or from one biological system to the other.

The enzymatic nucleic acid-based inhibitors of PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV expression are useful for the prevention of the diseases and conditions including HBV infection, hepatitis, hepatocellular carcinoma, tumorigenesis, cirrhosis, liver failure, cancers including breast, ovarian, prostate, and esophageal cancer, tumorigenesis, retinopathy, arthritis, psoriasis, female reproduction, restinosis, certain infectious diseases, transplant rejection and autoimmune disease such as multiple sclerosis, lupus, and AIDS, age related diseases such as macular degeneration and skin ulceration, Alzheimer's disease, dementia, diabetes, obesity and any other condition related to the level of PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV in a cell or tissue. and any other diseases or conditions that are related to the levels of PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV in a cell or tissue.

By "related" is meant that the reduction of PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV expression (specifically PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV genes) RNA levels and thus reduction in the level of the respective protein will relieve, to some extent, the symptoms of the disease or condition.

The nucleic acid-based inhibitors of the invention are added directly, or can be complexed with cationic lipids, packaged within liposomes, or otherwise delivered to target cells or tissues. The nucleic acid or nucleic acid complexes can be locally administered to relevant tissues *ex vivo*, or *in vivo* through injection, infusion pump or stent, with or without their incorporation in biopolymers. In preferred embodiments, the enzymatic nucleic acid inhibitors comprise sequences, which are complementary to the substrate sequences in Tables 3-31, 33, 34, 36-43, 56, 58, 59, 62, 63. Examples of such enzymatic nucleic acid molecules also are shown in Tables 3-29, 31, 33, 34, 37-43, 56, 58, 59, 62, 63. Examples of such enzymatic nucleic acid molecules consist essentially of sequences defined in these tables.

In yet another embodiment, the invention features antisense nucleic acid molecules including sequences complementary to the substrate sequences shown in Tables 3-31, 33, 34, 36, 37-43, 56, 58, 59, 62, 63. Such nucleic acid molecules can include sequences as shown for the binding arms of the enzymatic nucleic acid molecules in Tables 3-29, 31, 33, 34, 37-43, 56, 58, 59, 62, 63. Similarly, triplex molecules can be provided targeted to the corresponding DNA target regions, and containing the DNA equivalent of a target sequence or a sequence complementary to the specified target (substrate) sequence. Typically, antisense molecules will be complementary to a target sequence along a single contiguous sequence of the antisense molecule. However, in certain embodiments, an antisense molecule may bind to substrate such that the substrate molecule forms a loop, and/or an antisense molecule may bind such that the antisense molecule forms a loop. Thus, the antisense molecule may be complementary to two (or even more) non-contiguous substrate sequences or two (or even more) non-contiguous sequence portions of an antisense molecule may be complementary to a target sequence or both.

In another aspect, the invention provides mammalian cells containing one or more nucleic acid molecules and/or expression vectors of this invention. The one or more nucleic acid molecules may independently be targeted to the same or different sites.

By "consists essentially of" is meant that the active nucleic acid molecule of the invention, for example, an enzymatic nucleic acid molecule, contains an enzymatic center or core equivalent to those in the examples, and binding arms able to bind mRNA such that cleavage at the target site occurs. Other sequences may be present which do not interfere with such cleavage. Thus, a core region may, for example, include one or more loop or stem-loop structures, which do not prevent enzymatic activity. "X" in the sequences in Tables 3, 4, 9, 10, 13, 14, 18, 19, 24, 25, 33, 34, 37, 38, 63 can be such a loop. A core sequence for a hammerhead ribozyme can be CUGAUGAG X CGAA where X=GCCGUUAGGC or other stem II region as specifically or generally known in the art.

In another aspect of the invention, ribozymes or antisense molecules that interact with target RNA molecules and inhibit PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV (specifically PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV RNA) activity are expressed from transcription units inserted into DNA or RNA vectors. The recombinant vectors are preferably DNA plasmids or viral vectors. Ribozyme or antisense expressing viral vectors could be constructed based on, but not limited to, adeno-associated virus, retrovirus, adenovirus, or alphavirus. Preferably, the recombinant vectors capable of expressing the ribozymes or antisense are delivered as described above, and persist in target cells. Alternatively, viral vectors may be used that provide for transient expression of ribozymes or antisense. Such vectors might be repeatedly administered as necessary. Once expressed, the ribozymes or antisense bind to the target RNA and inhibit its function or expression. Delivery of ribozyme or antisense expressing vectors could be systemic, such as by intravenous or intramuscular administration, by administration to target cells ex-planted from the patient followed by reintroduction into the patient, or by any other means that would allow for introduction into the desired target cell. Antisense DNA can be expressed via the use of a single stranded DNA intracellular expression vector.

By RNA is meant a molecule comprising at least one ribonucleotide residue. By "ribonucleotide" is meant a nucleotide with a hydroxyl group at the 2' position of a β -D-ribo-furanose moiety.

By "vectors" is meant any nucleic acid- and/or viral-based technique used to deliver a desired nucleic acid.

By "patient" is meant an organism, which is a donor or recipient of explanted cells or the cells themselves. "Patient" also refers to an organism to which the nucleic acid molecules of the invention can be administered. Preferably, a patient is a mammal or mammalian cells. More preferably, a patient is a human or human cells.

5 The nucleic acid molecules of the instant invention, individually, or in combination or in conjunction with other drugs, can be used to treat diseases or conditions discussed above. For example, to treat a disease or condition associated with PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV, the patient may be treated, or other appropriate cells may be treated, as is evident to those skilled in the art, individually or in
10 combination with one or more drugs under conditions suitable for the treatment.

 In a further embodiment, the described molecules, such as antisense or ribozymes, can be used in combination with other known treatments to treat conditions or diseases discussed above. For example, the described molecules could be used in combination with one or more known therapeutic agents to treat HBV infection, hepatitis, hepatocellular
15 carcinoma, tumorigenesis, cirrhosis, liver failure, cancers including breast, ovarian, prostate, and esophageal cancer, tumorigenesis, retinopathy, arthritis, psoriasis, female reproduction, restinosis, certain infectious diseases, transplant rejection and autoimmune disease such as multiple sclerosis, lupus, and AIDS, age related diseases such as macular degeneration and skin ulceration, Alzheimer's disease, dementia, diabetes, and/or obesity.

20 In another preferred embodiment, the invention features nucleic acid-based inhibitors (*e.g.*, enzymatic nucleic acid molecules (ribozymes), antisense nucleic acids, triplex DNA, antisense nucleic acids containing RNA cleaving chemical groups) and methods for their use to down regulate or inhibit the expression of RNA (*e.g.*, PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV) capable of progression
25 and/or maintenance of HBV infection, hepatitis, hepatocellular carcinoma, tumorigenesis, cirrhosis, liver failure, cancers including breast, ovarian, prostate, and esophageal cancer, tumorigenesis, retinopathy, arthritis, psoriasis, female reproduction, restinosis, certain infectious diseases, transplant rejection and autoimmune disease such as multiple sclerosis, lupus, and AIDS, age related diseases such as macular degeneration and skin ulceration,
30 Alzheimer's disease, dementia, diabetes, and/or obesity.

In another preferred embodiment, the invention features nucleic acid-based techniques (*e.g.*, enzymatic nucleic acid molecules (ribozymes), antisense nucleic acids, triplex DNA, antisense nucleic acids containing RNA cleaving chemical groups) and methods for their use to down regulate or inhibit the expression of PTP-1B, MetAP-2,
5 BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV RNA expression.

By "comprising" is meant including, but not limited to, whatever follows the word "comprising". Thus, use of the term "comprising" indicates that the listed elements are required or mandatory, but that other elements are optional and may or may not be present. By "consisting of" is meant including, and limited to, whatever follows the phrase
10 "consisting of". Thus, the phrase "consisting of" indicates that the listed elements are required or mandatory, and that no other elements may be present. By "consisting essentially of" is meant including any elements listed after the phrase, and limited to other elements that do not interfere with or contribute to the activity or action specified in the disclosure for the listed elements. Thus, the phrase "consisting essentially of" indicates
15 that the listed elements are required or mandatory, but that other elements are optional and may or may not be present depending upon whether or not they affect the activity or action of the listed elements.

Other features and advantages of the invention will be apparent from the following description of the preferred embodiments thereof, and from the claims.

20

Description Of The Preferred Embodiments

The drawings will first briefly be described.

Drawings:

25 **Figure 1** shows the secondary structure model for seven different classes of enzymatic nucleic acid molecules. Arrow indicates the site of cleavage. ----- indicate the target sequence. Lines interspersed with dots are meant to indicate tertiary interactions. - is meant to indicate base-paired interaction. **Group I Intron:** P1-P9.0 represent various stem-loop structures (Cech *et al.*, 1994, *Nature Struc. Bio.*, 1, 273). **RNase P (M1RNA):**
30 EGS represents external guide sequence (Forster *et al.*, 1990, *Science*, 249, 783; Pace *et al.*, 1990, *J. Biol. Chem.*, 265, 3587). **Group II Intron:** 5'SS means 5' splice site; 3'SS means 3'-splice site; IBS means intron binding site; EBS means exon binding site (Pyle *et*

al., 1994, *Biochemistry*, 33, 2716). **VS RNA**: I-VI are meant to indicate six stem-loop structures; shaded regions are meant to indicate tertiary interaction (Collins, International PCT Publication No. WO 96/19577). **HDV Ribozyme**: I-IV are meant to indicate four stem-loop structures (Been *et al.*, US Patent No. 5,625,047). **Hammerhead Ribozyme**: I-III are meant to indicate three stem-loop structures; stems I-III can be of any length and may be symmetrical or asymmetrical (Usman *et al.*, 1996, *Curr. Op. Struct. Bio.*, 1, 527). **Hairpin Ribozyme**: Helix 1, 4 and 5 can be of any length; Helix 2 is between 3 and 8 base-pairs long; Y is a pyrimidine; Helix 2 (H2) is provided with a least 4 base pairs (*i.e.*, n is 1, 2, 3 or 4) and helix 5 can be optionally provided of length 2 or more bases (preferably 3 - 20 bases, *i.e.*, m is from 1 - 20 or more). Helix 2 and helix 5 may be covalently linked by one or more bases (*i.e.*, r is ≥ 1 base). Helix 1, 4 or 5 may also be extended by 2 or more base pairs (*e.g.*, 4 - 20 base pairs) to stabilize the ribozyme structure, and preferably is a protein binding site. In each instance, each N and N' independently is any normal or modified base and each dash represents a potential base-pairing interaction. These nucleotides may be modified at the sugar, base or phosphate. Complete base-pairing is not required in the helices, but is preferred. Helix 1 and 4 can be of any size (*i.e.*, o and p is each independently from 0 to any number, *e.g.*, 20) as long as some base-pairing is maintained. Essential bases are shown as specific bases in the structure, but those in the art will recognize that one or more may be modified chemically (abasic, base, sugar and/or phosphate modifications) or replaced with another base without significant effect. Helix 4 can be formed from two separate molecules, *i.e.*, without a connecting loop. The connecting loop when present may be a ribonucleotide with or without modifications to its base, sugar or phosphate. "q" \geq 2 bases. The connecting loop can also be replaced with a non-nucleotide linker molecule. H refers to bases A, U, or C. Y refers to pyrimidine bases. "_____" refers to a covalent bond. (Burke *et al.*, 1996, *Nucleic Acids & Mol. Biol.*, 10, 129; Chowrira *et al.*, US Patent No. 5,631,359).

Figure 2 shows examples of chemically stabilized ribozyme motifs. **HH Rz**, represents hammerhead ribozyme motif (Usman *et al.*, 1996, *Curr. Op. Struct. Bio.*, 1, 527); **NCH Rz** represents the NCH ribozyme motif (described herein and in Ludwig & Sproat, International PCT Publication No. WO 98/58058); **G-Cleaver**, represents G-cleaver ribozyme motif (Kore *et al.*, 1998, *Nucleic Acids Research*, 26, 4116-4120). N or

n, represent independently a nucleotide which may be same or different and have complementarity to each other; **rI**, represents ribo-Inosine nucleotide; arrow indicates the site of cleavage within the target. Position 4 of the HH Rz and the NCH Rz is shown as having 2'-C-allyl modification, but those skilled in the art will recognize that this position
5 can be modified with other modifications well known in the art, so long as such modifications do not significantly inhibit the activity of the ribozyme.

Figure 3 shows an example of the Amberzyme ribozyme motif that is chemically stabilized (see, for example, Beigelman *et al.*, International PCT publication No. WO 99/55857; also referred to as Class I Motif). The Amberzyme motif is a class of enzymatic
10 nucleic acid molecules that do not require the presence of a ribonucleotide (2'-OH) group for activity.

Figure 4 shows an example of the Zinzyme A ribozyme motif that is chemically stabilized (see, for example, International PCT publication No. WO 99/55857; also referred to as Class A Motif). The Zinzyme motif is a class of enzymatic nucleic acid
15 molecules that do not require the presence of a ribonucleotide (2'-OH) group for activity.

Figure 5 shows an example of a DNAzyme motif described by Santoro *et al.*, 1997, PNAS, 94, 4262.

Figure 6 is a diagrammatic representation of the hammerhead ribozyme motif known in the art and the NCH motif. Stem II can be 2 base-pair long, preferably, 2, 3,
20 4, 5, 6, 7, 8, and 10 base-pairs long. Each N and N' is independently any base or non-nucleotide as used herein; X is adenosine, cytidine or uridine; Stem I-III are meant to indicate three stem-loop structures; stems I-III can be of any length and may be symmetrical or asymmetrical (Usman *et al.*, 1996, *Curr. Op. Struct. Bio.*, 1, 527); arrow indicates the site of cleavage in the target RNA; Rz refers to ribozyme; Loop II may be
25 present or absent. If Loop II is present it is greater than or equal to three nucleotides, preferably four nucleotides. The Loop II sequence is preferably 5'-GAAA-3' or 5'-GUUA-3'.

Figure 7 shows examples of chemically stabilized ribozyme motifs. **HH Rz**, represents hammerhead ribozyme motif (Usman *et al.*, 1996, *Curr. Op. Struct. Bio.*, 1, 527); **NCH-Inosine Rz** represents the NCH ribozyme motif with riboinosine at 15.1
30 position; **NCH-Xylo Rz** represents the NCH ribozyme with xylo inosine at 15.1 position. N or n, represent independently a nucleotide which may be same or different and may have

complementarity to each other; **rI**, represents ribo-Inosine nucleotide; **xI** represent xylo-
inosine; arrow indicates the site of cleavage within the target. Position 4 of the HH RZ and
the NCH Rzs is shown as having 2'-C-allyl modification, but those skilled in the art will
recognize that this position can be modified with other modifications well known in the
art, so long as such modifications do not significantly inhibit the activity of the ribozyme.

Figure 8 is a graphical representation of data showing inhibition of cell proliferation
mediated by NCH and HH ribozymes targeted against HER2/*neu*/ErbB2 gene. Untreated,
refers to cells not treated with ribozymes; HH RZ refers to hammerhead ribozyme; NCX
RZ refers to the NCH ribozymes of the invention; IA refers to catalytically inactive or
attenuated ribozyme used as a control.

Figure 9 is a schematic diagram of the process for the synthesis of beta-D-
xylofuranosyl hypoxanthine 3'-phosphoramidite.

Figure 10 displays a schematic representation of NTP synthesis using nucleoside
substrates.

Figure 11 shows a scheme for an in vitro selection method. A pool of nucleic acid
molecules is generated with a random core region and one or more region(s) with a defined
sequence. These nucleic acid molecules are bound to a column containing immobilized
oligonucleotide with a defined sequence, where the defined sequence is complementary to
region(s) of defined sequence of nucleic acid molecules in the pool. Those nucleic acid
molecules capable of cleaving the immobilized oligonucleotide (target) in the column are
isolated and converted to complementary DNA (cDNA), followed by transcription using
NTPs to form a new nucleic acid pool.

Figure 12 shows a scheme for a two column in vitro selection method. A pool of
nucleic acid molecules is generated with a random core and two flanking regions (region A
and region B) with defined sequences. The pool is passed through a column which has
immobilized oligonucleotides with regions A' and B' that are complementary to regions A
and B of the nucleic acid molecules in the pool, respectively. The column is subjected to
conditions sufficient to facilitate cleavage of the immobilized oligonucleotide target. The
molecules in the pool that cleave the target (active molecules) have A' region of the target
bound to their A region, whereas the B region is free. The column is washed to isolate the
active molecules with the bound A' region of the target. This pool of active molecules
may also contain some molecules that are not active to cleave the target (inactive

molecules) but have dissociated from the column. To separate the contaminating inactive molecules from the active molecules, the pool is passed through a second column (column 2) which contains immobilized oligonucleotides with the A' sequence but not the B' sequence. The inactive molecules will bind to column 2 but the active molecules will not
5 bind to column 2 because their A region is occupied by the A' region of the target oligonucleotide from column 1. Column 2 is washed to isolate the active molecules for further processing as described in the scheme shown in **Figure 11**.

Figure 13 is a diagram of a novel 48 nucleotide enzymatic nucleic acid motif which was identified using in vitro methods described in the instant invention. The molecule
10 shown is only exemplary. The 5' and 3' terminal nucleotides (referring to the nucleotides of the substrate binding arms rather than merely the single terminal nucleotide on the 5' and 3' ends) can be varied so long as those portions can base-pair with target substrate sequence. In addition, the guanosine (G) shown at the cleavage site of the substrate can be changed to other nucleotides so long as the change does not eliminate the ability of
15 enzymatic nucleic acid molecules to cleave the target sequence. Substitutions in the nucleic acid molecule and/or in the substrate sequence can be readily tested, for example, as described herein.

Figure 14 is a schematic diagram of HCV luciferase assay used to demonstrate efficacy of class I enzymatic nucleic acid molecule motif.

20 **Figure 15** is a graph indicating the dose curve of an enzymatic nucleic acid molecule targeting site 146 on HCV RNA.

Figure 16 is a bar graph showing enzymatic nucleic acid molecules targeting 4 sites within the HCV RNA are able to reduce RNA levels in cells.

25 **Figure 17** shows secondary structures and cleavage rates for characterized Class II enzymatic nucleic acid motifs.

Figure 18 is a diagram of a novel 35 nucleotide enzymatic nucleic acid motif which was identified using in vitro methods described in the instant invention. The molecule shown is only exemplary. The 5' and 3' terminal nucleotides (referring to the nucleotides of the substrate binding arms rather than merely the single terminal nucleotide on the 5'
30 and 3' ends) can be varied so long as those portions can base-pair with target substrate sequence. In addition, the guanosine (G) shown at the cleavage site of the substrate can be changed to other nucleotides so long as the change does not eliminate the ability of

enzymatic nucleic acid molecules to cleave the target sequence. Substitutions in the nucleic acid molecule and/or in the substrate sequence can be readily tested, for example, as described herein.

5 **Figure 19** is a bar graph showing substrate specificities for Class II (zinzyme) ribozymes.

Figure 20 is a bar graph showing Class II enzymatic nucleic acid molecules targeting 10 representative sites within the HER2 RNA in a cellular proliferation screen.

Figure 21 is a synthetic scheme outlining the synthesis of 5-[3-aminopropynyl(propyl)]uridine 5'-triphosphates and 4-imidazoleaceticacid conjugates.

10 **Figure 22** is a synthetic scheme outlining the synthesis of 5-[3-(N-4-imidazoleacetyl)aminopropynyl(propyl)]uridine 5'-triphosphates.

Figure 23 is a synthetic scheme outlining the synthesis of carboxylate tethered uridine 5'-triphosphates.

15 **Figure 24** is a synthetic scheme outlining the synthesis of 5-(3-aminoalkyl) and 5-[3(N-succinyl)aminopropyl] functionalized cytidines.

Figure 25 is a diagram of a class I ribozyme stem truncation and loop replacement analysis.

Figure 26 is a diagram of class I ribozymes with truncated stem(s) and/or non-nucleotide linkers used in loop structures.

20 **Figure 27** is a diagram of "no-ribo" class II ribozymes.

Figure 28 is a graph showing cleavage reactions with class II ribozymes under differing divalent metal concentrations.

Figure 29 is a diagram of differing class II ribozymes with varying ribo content and their relative rates of catalysis.

25 **Figure 30** is a graph showing class II ribozyme (zinzyme) mediated reduction of HER2 RNA in SKBR3 breast carcinoma cells. Cells were treated with 100 nm, and 200 nm of zinzyme (RPI 18656) targeting site 972 of HER2 RNA and a corresponding scrambled attenuated control complexed with 2.5 µg/ml of lipid. Active zinzymes and scrambled attenuated controls were compared to untreated cells after 24 hours post
30 treatment.

Figure 31 is a graph showing class II ribozyme (zinzyme) mediated dose response anti-proliferation assay in SKBR3 breast carcinoma cells. Cells were treated with 100 nm, and 200 nm of zinzyme (RPI 18656) targeting site 972 of HER2 RNA and a corresponding scrambled attenuated control complexed with 2.0 µg/ml of lipid. Active zinzymes and scrambled attenuated controls were compared to untreated cells after 24 hours post treatment.

Figure 32 is a graph which shows the dose dependent reduction of HER2 RNA in SKOV-3 cells treated with RPI 19293 from 0 to 100 nM with 5.0 µg/ml of cationic lipid.

Figure 33 is a graph which shows the dose dependent reduction of HER2 RNA and inhibition of cellular proliferation in SKBR-3 cells treated with RPI 19293 from 0 to 400 nM with 5.0 µg/ml of cationic lipid.

Figure 34 shows a non-limiting example of the replacement of a 2'-O-methyl 5'-CA-3' with a ribo G in the class II (zinzyme) motif. The representative motif shown for the purpose of the figure is a "seven-ribo" zinzyme motif, however, the interchangeability of a G and a CA in the position shown in **Figure 25** of the class II (zinzyme) motif extends to any combination of 2-O-methyl and ribo residues. For instance, a 2'-O-methyl G can replace the 2'-O-methyl 5'-CA-3' and vice versa.

Figure 35 is a graph which shows a screen of class II ribozymes (zinzymes) targeting site 972 of HER2 RNA which contain ribo-G reductions (RPI 19727 = no ribo, RPI 19728 = one ribo, RPI 19293 = two ribo, RPI 19729 = three ribo, RPI 19730 = four ribo, 19731 = five ribo, and RPI 19292 = seven ribo) for anti-proliferative activity in SKBR3 cells.

Figure 36 summarizes the results of functional group modification studies in which various nucleoside analogs were tested for activity in the NCH ribozyme motif. K_{rel} values describe the cleavage values of a given substituent at position 15.1 relative the Inosine at position 15.1 (I-15.1).

Figure 37 summarizes reported functional group modification studies performed at the A 15.1 residue in the A-15.1 •U-16.1 context of NUH cleaving ribozymes. K_{rel} values describe the cleavage values of a given substituent at position 15.1 relative the adenosine at position 15.1 (A-15.1).

Mechanism of action of Nucleic Acid Molecules of the Invention

Antisense: Antisense molecules may be modified or unmodified RNA, DNA, or mixed polymer oligonucleotides and primarily function by specifically binding to matching sequences resulting in inhibition of peptide synthesis (Wu-Pong, Nov 1994, *BioPharm*, 20-33). The antisense oligonucleotide binds to target RNA by Watson Crick base-pairing and blocks gene expression by preventing ribosomal translation of the bound sequences either by steric blocking or by activating RNase H enzyme. Antisense molecules may also alter protein synthesis by interfering with RNA processing or transport from the nucleus into the cytoplasm (Mukhopadhyay & Roth, 1996, *Crit. Rev. in Oncogenesis* 7, 151-190).

In addition, binding of single stranded DNA to RNA may result in nuclease degradation of the heteroduplex (Wu-Pong, *supra*; Crooke, *supra*). To date, the only backbone modified DNA chemistry which will act as substrates for RNase H are phosphorothioates, phosphorodithioates, and borontrifluoridates. Recently, it has been reported that 2'-arabino and 2'-fluoro arabino- containing oligos can also activate RNase H activity.

A number of antisense molecules have been described that utilize novel configurations of chemically modified nucleotides, secondary structure, and/or RNase H substrate domains (Woelf *et al.*, International PCT Publication No. WO 98/13526; Thompson *et al.*, International PCT Publication No. WO 99/54459 ; Hartmann *et al.*, International PCT Publication No. WO 00/17346) all of these are incorporated by reference herein in their entirety.

Antisense DNA can be used to target RNA by means of DNA-RNA interactions, thereby activating RNase H, which digests the target RNA in the duplex. Antisense DNA can be chemically synthesized or can be expressed via the use of a single stranded DNA intracellular expression vector or the equivalent thereof.

Triplex Forming Oligonucleotides (TFO): Single stranded DNA may be designed to bind to genomic DNA in a sequence specific manner. TFOs are comprised of pyrimidine-rich oligonucleotides which bind DNA helices through Hoogsteen Base-pairing (Wu-Pong, *supra*). The resulting triple helix composed of the DNA sense, DNA antisense, and TFO disrupts RNA synthesis by RNA polymerase. The TFO mechanism may result in gene expression or cell death since binding may be irreversible (Mukhopadhyay & Roth, *supra*)

2'-5' Oligoadenylates: The 2-5 A system is an interferon-mediated mechanism for RNA degradation found in higher vertebrates (Mitra *et al.*, 1996, *Proc Nat Acad Sci USA* 93, 6780-6785). Two types of enzymes, 2-5A synthetase and RNase L, are required for RNA cleavage. The 2-5A synthetases require double stranded RNA to form 2'-5' oligoadenylates (2-5A). 2-5A then acts as an allosteric effector for utilizing RNase L which has the ability to cleave single stranded RNA. The ability to form 2-5A structures with double stranded RNA makes this system particularly useful for inhibition of viral replication.

(2'-5') oligoadenylate structures may be covalently linked to antisense molecules to form chimeric oligonucleotides capable of RNA cleavage (Torrence, *supra*). These molecules putatively bind and activate a 2-5A dependent RNase, the oligonucleotide/enzyme complex then binds to a target RNA molecule which can then be cleaved by the RNase enzyme. The covalent attachment of 2'-5' oligoadenylate structures is not limited to antisense applications, and can be further elaborated to include attachment to nucleic acid molecules of the instant invention.

Enzymatic Nucleic Acid: Seven basic varieties of naturally-occurring enzymatic RNAs are presently known. In addition, several *in vitro* selection (evolution) strategies (Orgel, 1979, *Proc. R. Soc. London, B* 205, 435) have been used to evolve new nucleic acid catalysts capable of catalyzing cleavage and ligation of phosphodiester linkages (Joyce, 1989, *Gene*, 82, 83-87; Beaudry *et al.*, 1992, *Science* 257, 635-641; Joyce, 1992, *Scientific American* 267, 90-97; Breaker *et al.*, 1994, *TIBTECH* 12, 268; Bartel *et al.*, 1993, *Science* 261:1411-1418; Szostak, 1993, *TIBS* 17, 89-93; Kumar *et al.*, 1995, *FASEB J.*, 9, 1183; Breaker, 1996, *Curr. Op. Biotech.*, 7, 442; Santoro *et al.*, 1997, *Proc. Natl. Acad. Sci.*, 94, 4262; Tang *et al.*, 1997, *RNA* 3, 914; Nakamaye & Eckstein, 1994, *supra*; Long & Uhlenbeck, 1994, *supra*; Ishizaka *et al.*, 1995, *supra*; Vaish *et al.*, 1997, *Biochemistry* 36, 6495; all of these are incorporated by reference herein). Each can catalyze a series of reactions including the hydrolysis of phosphodiester bonds in *trans* (and thus can cleave other RNA molecules) under physiological conditions.

In general, enzymatic nucleic acids act by first binding to a target RNA. Such binding occurs through the target binding portion of an enzymatic nucleic acid which is held in close proximity to an enzymatic portion of the molecule that acts to cleave the target RNA. Thus, the enzymatic nucleic acid first recognizes and then binds a target

RNA through complementary base-pairing, and once bound to the correct site, acts enzymatically to cut the target RNA. Strategic cleavage of such a target RNA will destroy its ability to direct synthesis of an encoded protein. After an enzymatic nucleic acid has bound and cleaved its RNA target, it is released from that RNA to search for another target and can repeatedly bind and cleave new targets.

Nucleic acid molecules of this invention will block to some extent PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV protein expression and can be used to treat disease or diagnose disease associated with the levels of PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV.

The enzymatic nature of a ribozyme has significant advantages, such as the concentration of ribozyme necessary to affect a therapeutic treatment is low. This advantage reflects the ability of the ribozyme to act enzymatically. Thus, a single ribozyme molecule is able to cleave many molecules of target RNA. In addition, the ribozyme is a highly specific inhibitor, with the specificity of inhibition depending not only on the base-pairing mechanism of binding to the target RNA, but also on the mechanism of target RNA cleavage. Single mismatches, or base-substitutions, near the site of cleavage can be chosen to completely eliminate catalytic activity of a ribozyme.

Nucleic acid molecules having an endonuclease enzymatic activity are able to repeatedly cleave other separate RNA molecules in a nucleotide base sequence-specific manner. Such enzymatic nucleic acid molecules can be targeted to virtually any RNA transcript, and achieve efficient cleavage *in vitro* (Zaug *et al.*, 324, *Nature*, 429 1986 ; Uhlenbeck, 1987 *Nature*, 328, 596; Kim *et al.*, 84 *Proc. Natl. Acad. Sci. USA*, 8788, 1987; Dreyfus, 1988, *Einstein Quart. J. Bio. Med.*, 6, 92; Haseloff and Gerlach, 334 *Nature*, 585, 1988; Cech, 260 *JAMA*, 3030, 1988; Jefferies *et al.*, 17 *Nucleic Acids Research*, 1371, 1989; and Santoro *et al.*, 1997 *supra*).

Because of their sequence specificity, *trans*-cleaving ribozymes show promise as therapeutic agents for human disease (Usman & McSwiggen, 1995 *Ann. Rep. Med. Chem.* 30, 285-294; Christoffersen and Marr, 1995 *J. Med. Chem.* 38, 2023-2037). Ribozymes can be designed to cleave specific RNA targets within the background of cellular RNA. Such a cleavage event renders the RNA non-functional and abrogates protein expression from that RNA. In this manner, synthesis of a protein associated with a disease state can be selectively inhibited (Warashina *et al.*, 1999, *Chemistry and Biology*, 6, 237-250).

The nucleic acid molecules of the instant invention are also referred to as GeneBloc™ reagents, which are essentially nucleic acid molecules (e.g.; ribozymes, antisense) capable of down-regulating gene expression.

5 Target sites

Targets for useful ribozymes and antisense nucleic acids can be determined as disclosed in Draper *et al.*, WO 93/23569; Sullivan *et al.*, WO 93/23057; Thompson *et al.*, WO 94/02595; Draper *et al.*, WO 95/04818; McSwiggen *et al.*, US Patent No. 5,525,468, and all hereby incorporated in their entireties by reference herein. Other examples include

10 the following PCT applications, which concern inactivation of expression of disease-related genes: WO 95/23225, WO 95/13380, WO 94/02595, all incorporated by reference herein. Rather than repeat the guidance provided in those documents here, below are provided specific examples of such methods, not limiting to those in the art. Ribozymes and antisense to such targets are designed as described in those applications and

15 synthesized to be tested *in vitro* and *in vivo*, as also described. The sequence of human PTP-1B, MetAP-2, BACE, ps-1, ps-2, HER2, PLN, TERT, and/or HBV RNAs (for example, GenBank accession Nos. (PTP-1B, NM_002827), (MetAP-2, U29607), (BACE, AF190725), (ps-1, L76517), (ps-2, L43964), (HER2/c-erb2/neu, X03363), (PLN, NM_002667), (TERT, NM_003219) and (HBV, AF100308.1, HBV strain 2-18;

20 additionally, other HBV strains can be screened by one skilled in the art, see Table 35 for other possible strains) were screened for optimal enzymatic nucleic acid and antisense target sites using a computer-folding algorithm. Antisense, hammerhead, DNAzyme, NCH (Inozyme), amberzyme, zinzyme or G-Cleaver ribozyme binding/cleavage sites were identified. These sites are shown in Tables 3-29, 31, 33, 34, 37-43, 56, 58, 59, 62, 63 (all

25 sequences are 5' to 3' in the tables; X can be any base-paired sequence, the actual sequence is not relevant here). The nucleotide base position is noted in the Tables as that site to be cleaved by the designated type of enzymatic nucleic acid molecule. Table 36 shows substrate positions selected from Renbo *et al.*, 1987, *Sci. Sin.*, 30, 507, used in Draper, US patent No. 6,017,756 entitled "METHOD AND REAGENT FOR

30 INHIBITING HEPATITIS B VIRUS REPLICATION" and Draper *et al.*, International PCT publication No. WO 93/23569, filed April 29, 1993, entitled "METHOD AND REAGENT FOR INHIBITING VIRAL REPLICATION". While human sequences can be

screened and enzymatic nucleic acid molecule and/or antisense thereafter designed, as discussed in Stinchcomb *et al.*, WO 95/23225, mouse targeted ribozymes may be useful to test efficacy of action of the enzymatic nucleic acid molecule and/or antisense prior to testing in humans.

5 Antisense, hammerhead, DNAzyme, NCH (Inozyme), amberzyme, zinzyme or G-Cleaver ribozyme binding/cleavage sites were identified, as discussed above. The nucleic acid molecules were individually analyzed by computer folding (Jaeger *et al.*, 1989 *Proc. Natl. Acad. Sci. USA*, 86, 7706) to assess whether the sequences fold into the appropriate secondary structure. Those nucleic acid molecules with unfavorable intramolecular
10 interactions such as between the binding arms and the catalytic core were eliminated from consideration. Varying binding arm lengths can be chosen to optimize activity.

 Antisense, hammerhead, DNAzyme, NCH, amberzyme, zinzyme or G-Cleaver ribozyme binding/cleavage sites were identified and were designed to anneal to various sites in the RNA target. The binding arms are complementary to the target site sequences
15 described above. The nucleic acid molecules were chemically synthesized. The method of synthesis used follows the procedure for normal DNA/RNA synthesis as described below and in Usman *et al.*, 1987 *J. Am. Chem. Soc.*, 109, 7845; Scaringe *et al.*, 1990 *Nucleic Acids Res.*, 18, 5433; Wincott *et al.*, 1995 *Nucleic Acids Res.* 23, 2677-2684; and Caruthers *et al.*, 1992, *Methods in Enzymology* 211,3-19.

20

Synthesis of Nucleic acid Molecules

 Synthesis of nucleic acids greater than 100 nucleotides in length is difficult using automated methods, and the therapeutic cost of such molecules is prohibitive. In this invention, small nucleic acid motifs ("small refers to nucleic acid motifs no more than 100
25 nucleotides in length, preferably no more than 80 nucleotides in length, and most preferably no more than 50 nucleotides in length; *e.g.*, antisense oligonucleotides, hammerhead or the NCH ribozymes) are preferably used for exogenous delivery. The simple structure of these molecules increases the ability of the nucleic acid to invade targeted regions of RNA structure. Exemplary molecules of the instant invention are
30 chemically synthesized, and others can similarly be synthesized.

Oligonucleotides (e.g.; antisense GeneBlocs) are synthesized using protocols known in the art as described in Caruthers *et al.*, 1992, *Methods in Enzymology* 211, 3-19, Thompson *et al.*, International PCT Publication No. WO 99/54459, Wincott *et al.*, 1995, *Nucleic Acids Res.* 23, 2677-2684, Wincott *et al.*, 1997, *Methods Mol. Bio.*, 74, 59, Brennan *et al.*, 1998, *Biotechnol Bioeng.*, 61, 33-45, and Brennan, US patent No. 6,001,311. All of these references are incorporated herein by reference. The synthesis of oligonucleotides makes use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. In a non-limiting example, small scale syntheses are conducted on a 394 Applied Biosystems, Inc. synthesizer using a 0.2 μ mol scale protocol with a 2.5 min coupling step for 2'-O-methylated nucleotides and a 45 sec coupling step for 2'-deoxy nucleotides. Table II outlines the amounts and the contact times of the reagents used in the synthesis cycle. Alternatively, syntheses at the 0.2 μ mol scale can be performed on a 96-well plate synthesizer, such as the instrument produced by Protogene (Palo Alto, CA) with minimal modification to the cycle. A 33-fold excess (60 μ L of 0.11 M = 6.6 μ mol) of 2'-O-methyl phosphoramidite and a 105-fold excess of S-ethyl tetrazole (60 μ L of 0.25 M = 15 μ mol) can be used in each coupling cycle of 2'-O-methyl residues relative to polymer-bound 5'-hydroxyl. A 22-fold excess (40 μ L of 0.11 M = 4.4 μ mol) of deoxy phosphoramidite and a 70-fold excess of S-ethyl tetrazole (40 μ L of 0.25 M = 10 μ mol) can be used in each coupling cycle of deoxy residues relative to polymer-bound 5'-hydroxyl. Average coupling yields on the 394 Applied Biosystems, Inc. synthesizer, determined by colorimetric quantitation of the trityl fractions, are typically 97.5-99%. Other oligonucleotide synthesis reagents for the 394 Applied Biosystems, Inc. synthesizer include the following: detritylation solution is 3% TCA in methylene chloride (ABI); capping is performed with 16% *N*-methyl imidazole in THF (ABI) and 10% acetic anhydride/10% 2,6-lutidine in THF (ABI); and oxidation solution is 16.9 mM I₂, 49 mM pyridine, 9% water in THF (PERSEPTIVE™). Burdick & Jackson Synthesis Grade acetonitrile is used directly from the reagent bottle. S-Ethyltetrazole solution (0.25 M in acetonitrile) is made up from the solid obtained from American International Chemical, Inc. Alternately, for the introduction of phosphorothioate linkages, Beaucage reagent (3H-1,2-Benzodithiol-3-one 1,1-dioxide, 0.05 M in acetonitrile) is used.

Deprotection of the antisense oligonucleotides is performed as follows: the polymer-bound trityl-on oligoribonucleotide is transferred to a 4 mL glass screw top vial and suspended in a solution of 40% aq. methylamine (1 mL) at 65 °C for 10 min. After cooling to -20 °C, the supernatant is removed from the polymer support. The support is
5 washed three times with 1.0 mL of EtOH:MeCN:H₂O/3:1:1, vortexed and the supernatant is then added to the first supernatant. The combined supernatants, containing the oligoribonucleotide, are dried to a white powder.

The method of synthesis used for normal RNA including certain enzymatic nucleic acid molecules follows the procedure as described in Usman *et al.*, 1987, *J. Am. Chem. Soc.*, 109, 7845; Scaringe *et al.*, 1990, *Nucleic Acids Res.*, 18, 5433; and Wincott *et al.*,
10 1995, *Nucleic Acids Res.* 23, 2677-2684 Wincott *et al.*, 1997, *Methods Mol. Bio.*, 74, 59, and makes use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. In a non-limiting example, small scale syntheses are conducted on a 394 Applied Biosystems, Inc.
15 synthesizer using a 0.2 µmol scale protocol with a 7.5 min coupling step for alkylsilyl protected nucleotides and a 2.5 min coupling step for 2'-O-methylated nucleotides. Table II outlines the amounts and the contact times of the reagents used in the synthesis cycle. Alternatively, syntheses at the 0.2 µmol scale can be done on a 96-well plate synthesizer, such as the instrument produced by Protogene (Palo Alto, CA) with minimal modification
20 to the cycle. A 33-fold excess (60 µL of 0.11 M = 6.6 µmol) of 2'-O-methyl phosphoramidite and a 75-fold excess of S-ethyl tetrazole (60 µL of 0.25 M = 15 µmol) can be used in each coupling cycle of 2'-O-methyl residues relative to polymer-bound 5'-hydroxyl. A 66-fold excess (120 µL of 0.11 M = 13.2 µmol) of alkylsilyl (ribo) protected phosphoramidite and a 150-fold excess of S-ethyl tetrazole (120 µL of 0.25 M = 30 µmol)
25 can be used in each coupling cycle of ribo residues relative to polymer-bound 5'-hydroxyl. Average coupling yields on the 394 Applied Biosystems, Inc. synthesizer, determined by colorimetric quantitation of the trityl fractions, are typically 97.5-99%. Other oligonucleotide synthesis reagents for the 394 Applied Biosystems, Inc. synthesizer include the following: detritylation solution is 3% TCA in methylene chloride (ABI);
30 capping is performed with 16% *N*-methyl imidazole in THF (ABI) and 10% acetic anhydride/10% 2,6-lutidine in THF (ABI); oxidation solution is 16.9 mM I₂, 49 mM pyridine, 9% water in THF (PERSEPTIVE™). Burdick & Jackson Synthesis Grade

acetonitrile is used directly from the reagent bottle. S-Ethyltetrazole solution (0.25 M in acetonitrile) is made up from the solid obtained from American International Chemical, Inc. Alternately, for the introduction of phosphorothioate linkages, Beaucage reagent (3H-1,2-Benzodithiol-3-one 1,1-dioxide 0.05 M in acetonitrile) is used.

5 Deprotection of the RNA is performed using either a two-pot or one-pot protocol. For the two-pot protocol, the polymer-bound trityl-on oligoribonucleotide is transferred to a 4 mL glass screw top vial and suspended in a solution of 40% aq. methylamine (1 mL) at 65 °C for 10 min. After cooling to -20 °C, the supernatant is removed from the polymer support. The support is washed three times with 1.0 mL of EtOH:MeCN:H₂O/3:1:1,
10 vortexed and the supernatant is then added to the first supernatant. The combined supernatants, containing the oligoribonucleotide, are dried to a white powder. The base deprotected oligoribonucleotide is resuspended in anhydrous TEA/HF/NMP solution (300 µL of a solution of 1.5 mL N-methylpyrrolidinone, 750 µL TEA and 1 mL TEA•3HF to provide a 1.4 M HF concentration) and heated to 65 °C. After 1.5 h, the oligomer is
15 quenched with 1.5 M NH₄HCO₃.

 Alternatively, for the one-pot protocol, the polymer-bound trityl-on oligoribonucleotide is transferred to a 4 mL glass screw top vial and suspended in a solution of 33% ethanolic methylamine/DMSO: 1/1 (0.8 mL) at 65 °C for 15 min. The vial is brought to r.t. TEA•3HF (0.1 mL) is added and the vial is heated at 65 °C for 15
20 min. The sample is cooled at -20 °C and then quenched with 1.5 M NH₄HCO₃.

 For purification of the trityl-on oligomers, the quenched NH₄HCO₃ solution is loaded onto a C-18 containing cartridge that had been prewashed with acetonitrile followed by 50 mM TEAA. After washing the loaded cartridge with water, the RNA is detritylated with 0.5% TFA for 13 min. The cartridge is then washed again with water,
25 salt exchanged with 1 M NaCl and washed with water again. The oligonucleotide is then eluted with 30% acetonitrile.

 Inactive hammerhead ribozymes or binding attenuated control (BAC) oligonucleotides) are synthesized by substituting a U for G₅ and a U for A₁₄ (numbering from Hertel, K. J., *et al.*, 1992, *Nucleic Acids Res.*, 20, 3252). Similarly, one or more
30 nucleotide substitutions can be introduced in other enzymatic nucleic acid molecules to inactivate the molecule and such molecules can serve as a negative control.

The average stepwise coupling yields are typically >98% (Wincott *et al.*, 1995 *Nucleic Acids Res.* 23, 2677-2684). Those of ordinary skill in the art will recognize that the scale of synthesis can be adapted to be larger or smaller than the example described above including but not limited to 96-well format, all that is important is the ratio of
5 chemicals used in the reaction.

Alternatively, the nucleic acid molecules of the present invention can be synthesized separately and joined together post-synthetically, for example, by ligation (Moore *et al.*, 1992, *Science* 256, 9923; Draper *et al.*, International PCT publication No. WO 93/23569; Shabarova *et al.*, 1991, *Nucleic Acids Research* 19, 4247; Bellon *et al.*, 1997, *Nucleosides & Nucleotides*, 16, 951; Bellon *et al.*, 1997, *Bioconjugate Chem.* 8, 204).
10

The nucleic acid molecules of the present invention are modified extensively to enhance stability by modification with nuclease resistant groups, for example, 2'-amino, 2'-C-allyl, 2'-fluoro, 2'-O-methyl, 2'-H (for a review see Usman and Cedergren, 1992, *TIBS* 17, 34; Usman *et al.*, 1994, *Nucleic Acids Symp. Ser.* 31, 163). Ribozymes are purified by
15 gel electrophoresis using general methods or are purified by high pressure liquid chromatography (HPLC; see Wincott *et al.*, *supra*, the totality of which is hereby incorporated herein by reference) and are re-suspended in water.

The sequences of the ribozymes and antisense constructs that are chemically synthesized, useful in this study, are shown in Tables 3-31, 33, 34, 37-43, 56, 58, 59, 62,
20 63. Those in the art will recognize that these sequences are representative only of many more such sequences where the enzymatic portion of the ribozyme (all but the binding arms) is altered to affect activity. The ribozyme and antisense construct sequences listed in Tables 3-31, 33, 34, 37-43, 56, 58, 59, 62, 63 may be formed of ribonucleotides or other nucleotides or non-nucleotides. Such ribozymes with enzymatic activity are equivalent to
25 the ribozymes described specifically in the Tables.

Optimizing Activity of the nucleic acid molecule of the invention.

Chemically synthesizing nucleic acid molecules with modifications (base, sugar and/or phosphate) that prevent their degradation by serum ribonucleases may increase their potency (see *e.g.*, Eckstein *et al.*, International Publication No. WO 92/07065; Perrault *et al.*, 1990 *Nature* 344, 565; Pieken *et al.*, 1991, *Science* 253, 314; Usman and Cedergren, 1992, *Trends in Biochem. Sci.* 17, 334; Usman *et al.*, International Publication No. WO
30 93/15187; Rossi *et al.*, International Publication No. WO 91/03162; Sproat, US Patent No.

5,334,711; and Burgin *et al.*, *supra*; all of these describe various chemical modifications that can be made to the base, phosphate and/or sugar moieties of the nucleic acid molecules herein and are all hereby incorporated by reference herein). Modifications which enhance their efficacy in cells, and removal of bases from nucleic acid molecules to shorten oligonucleotide synthesis times and reduce chemical requirements are desired.

There are several examples in the art describing sugar, base and phosphate modifications that can be introduced into nucleic acid molecules (e.g., enzymatic nucleic acid molecules) without significantly effecting catalysis and with significant enhancement in their nuclease stability and efficacy. Enzymatic nucleic acid molecules are modified to enhance stability and/or enhance catalytic activity by modification with nuclease resistant groups, for example, 2'-amino, 2'-C-allyl, 2'-fluoro, 2'-O-methyl, 2'-O-allyl, 2'-H, nucleotide base modifications (for a review see Usman and Cedergren, 1992 *TIBS* 17, 34; Usman *et al.*, 1994 *Nucleic Acids Symp. Ser.* 31, 163; Burgin *et al.*, 1996 *Biochemistry* 35, 14090). Sugar modification of enzymatic nucleic acid molecules have been extensively described in the art (see Eckstein *et al.*, *International Publication* PCT No. WO 92/07065; Perrault *et al.* *Nature* 1990, 344, 565-568; Pieken *et al.* *Science* 1991, 253, 314-317; Usman and Cedergren, *Trends in Biochem. Sci.* 1992, 17, 334-339; Usman *et al.* *International Publication* PCT No. WO 93/15187; Sproat, *US Patent* No. 5,334,711 and Beigelman *et al.*, 1995 *J. Biol. Chem.* 270, 25702; all of the references are hereby incorporated in their totality by reference herein). Such publications describe general methods and strategies to determine the location of incorporation of sugar, base and/or phosphate modifications and the like into enzymatic nucleic acid molecules without inhibiting catalysis, and are incorporated by reference herein. The 2'-position of the sugar in a nucleotide present in the nucleic acid molecules of the instant invention which tolerates substitution is selected from the group comprising -H, -OH, -COOH, -CONH₂, -CONHR¹, -CONR¹R², -NH₂, -NHR¹, -NR¹R², -NHCOR¹, -SH, SR¹, -F, -ONH₂, -ONHR¹, -ONR¹R², -NHOH, -NHOR¹, -NR²OH, -NR²OR¹, substituted or unsubstituted C₁-C₁₀ straight chain or branched alkyl, substituted or unsubstituted C₂-C₁₀ straight chain or branched alkenyl, substituted or unsubstituted C₂-C₁₀ straight chain or branched alkynyl, substituted or unsubstituted C₁-C₁₀ straight chain or branched alkoxy, substituted or unsubstituted C₂-C₁₀ straight chain or branched alkenyloxy, and substituted or unsubstituted C₂-C₁₀ straight chain or branched alkynyloxy. The substituents for sugar 2'

position preferably are independently halogen, cyano, amino, carboxy, ester, ether, carboxamide, hydroxy, or mercapto. R^1 and R^2 can be substituted or unsubstituted alkyl, alkenyl, or alkynyl groups, where the substituents are independently halogen, cyano, amino, carboxy, ester, ether, carboxamide, hydroxy, or mercapto.

5 In view of such teachings, similar modifications can be used as described herein to modify the nucleic acid molecules of the instant invention. Such publications describe general methods and strategies to determine the location of incorporation of sugar, base and/or phosphate modifications and the like into ribozymes without inhibiting catalysis, and are incorporated by reference herein. In view of such teachings, similar modifications
10 can be used as described herein to modify the nucleic acid molecules of the instant invention.

Some of the non-limiting examples of base modifications that can be introduced into enzymatic nucleic acids without significantly effecting their catalytic activity include, inosine, purine, pyridin-4-one, pyridin-2-one, phenyl, pseudouracil, 2, 4, 6-trimethoxy
15 benzene, 3-methyluracil, dihydrouridine, naphthyl, aminophenyl, 5-alkylcytidines (*e.g.*, 5-methylcytidine), 5-alkyluridines (*e.g.*, ribothymidine), 5-halouridine (*e.g.*, 5-bromouridine) or 6-azapyrimidines or 6-alkylpyrimidines (*e.g.* 6-methyluridine) and others (Burgin *et al.*, 1996, *Biochemistry*, 35, 14090). By "modified bases" in this aspect is
20 meant nucleotide bases other than adenine, guanine, cytosine and uracil at 1' position or their equivalents; such bases may be used within the catalytic core of the enzyme and/or in the substrate-binding regions.

The nucleic acid bases can be hypoxanthin-9-yl, or a functional equivalent thereof, in position^{15.1} of the ribozyme; the base at other positions may be guanin-9-yl, hypoxanthin-9-yl or 7-deazaguanin-9-yl in positions 5, 8 and 12 in the ribozyme; adenin-9-
25 yl, 2,6-diaminopurin-9-yl, purin-9-yl or 7-deaza adenin-9-yl in positions 6, 9, 13 and 14; uracil-1-yl, uracil-5-yl, thymine-1-yl or 5-propynyluracil-1-yl in position 4; cytosin-1-yl, 5-methylcytosin-1-yl or 5-propynylcytosin-1-yl in position 3; and adenin-9-yl, cytosin-1-yl, guanin-9-yl, uracil-1-yl, uracil-5-yl, hypoxanthin-9-yl, thymine-1-yl, 5-methylcytosin-1-yl, 2,6-diaminopurin-9-yl, purin-9-yl, 7-deaza adenin-9-yl, 7-deazaguanin-9-yl, 5-
30 propynylcytosin-1-yl, 5-propynyluracil-1-yl, isoguanin-9-yl, 2-aminopurin-9-yl, 6-methyluracil-1-yl, 4-thiouracil-1-yl, 2-pyrimidone-1-yl, quinazoline-2,4-dione-1-yl, xanthin-9-yl, N²-dimethylguanin-9-yl, or a functional equivalent thereof in position 7. The

base at position 15.1 is preferably hypoxanthin-9-yl or an analog where no hydrogen bond can form between any group at the 2 position of the base and the 2-oxo group of C^{16.1}.

Preferably, B is not guanin-9-yl in position 15.1.

5 In particular, the invention features modified ribozymes having a base substitution selected from pyridin-4-one, pyridin-2-one, phenyl, pseudouracil, 2, 4, 6-trimethoxy benzene, 3-methyluracil, dihydrouracil, naphthyl, 6-methyl-uracil and aminophenyl.

While chemical modification of oligonucleotide internucleotide linkages with phosphorothioate, phosphorothioate, and/or 5'-methylphosphonate linkages improves stability, too many of these modifications may cause some toxicity. Therefore, when
10 designing nucleic acid molecules, the amount of these internucleotide linkages should be minimized. The reduction in the concentration of these linkages should lower toxicity resulting in increased efficacy and higher specificity of these molecules.

Nucleic acid molecules having chemical modifications which maintain or enhance activity are provided. Such nucleic acid molecules are also generally more resistant to
15 nucleases than unmodified nucleic acid. Thus, in a cell and/or *in vivo* the activity may not be significantly lowered. Therapeutic nucleic acid molecules delivered exogenously must optimally be stable within cells until translation of the target RNA has been inhibited long enough to reduce the levels of the undesirable protein. This period of time varies between hours to days depending upon the disease state. Clearly, nucleic acid molecules must be
20 resistant to nucleases in order to function as effective intracellular therapeutic agents. Improvements in the chemical synthesis of RNA and DNA (Wincott *et al.*, 1995 *Nucleic Acids Res.* 23, 2677; Caruthers *et al.*, 1992, *Methods in Enzymology* 211,3-19 (all are incorporated by reference herein) have expanded the ability to modify nucleic acid molecules by introducing nucleotide modifications to enhance their nuclease stability as
25 described above.

Use of these the nucleic acid-based molecules of the invention will lead to better treatment of the disease progression by affording the possibility of combination therapies (e.g., multiple antisense or enzymatic nucleic acid molecules targeted to different genes, nucleic acid molecules coupled with known small molecule inhibitors, or intermittent
30 treatment with combinations of molecules (including different motifs) and/or other chemical or biological molecules). The treatment of patients with nucleic acid molecules may also include combinations of different types of nucleic acid molecules.

Therapeutic nucleic acid molecules (*e.g.*, enzymatic nucleic acid molecules and antisense nucleic acid molecules) delivered exogenously must optimally be stable within cells until translation of the target RNA has been inhibited long enough to reduce the levels of the undesirable protein. This period of time varies between hours to days depending upon the disease state. Clearly, these nucleic acid molecules must be resistant to nucleases in order to function as effective intracellular therapeutic agents. Improvements in the chemical synthesis of nucleic acid molecules described in the instant invention and in the art have expanded the ability to modify nucleic acid molecules by introducing nucleotide modifications to enhance their nuclease stability as described above.

By "enhanced enzymatic activity" is meant to include activity measured in cells and/or *in vivo* where the activity is a reflection of both catalytic activity and ribozyme stability. In this invention, the product of these properties is increased or not significantly (less than 10-fold) decreased *in vivo* compared to an all RNA ribozyme or all DNA enzyme.

In yet another preferred embodiment, nucleic acid catalysts having chemical modifications which maintain or enhance enzymatic activity are provided. Such nucleic acid catalysts are also generally more resistant to nucleases than unmodified nucleic acid. Thus, in a cell and/or *in vivo* the activity may not be significantly lowered. As exemplified herein such ribozymes are useful in a cell and/or *in vivo* even if activity over all is reduced 10 fold (Burgin *et al.*, 1996, *Biochemistry*, 35, 14090). Such ribozymes herein are said to "maintain" the enzymatic activity of an all RNA ribozyme.

In another aspect the nucleic acid molecules comprise a 5' and/or a 3'-cap structure.

By "cap structure" is meant chemical modifications, which have been incorporated at either terminus of the oligonucleotide (see, for example, Wincott *et al.*, WO 97/26270, incorporated by reference herein). These terminal modifications protect the nucleic acid molecule from exonuclease degradation, and may help in delivery and/or localization within a cell. The cap may be present at the 5'-terminus (5'-cap) or at the 3'-terminal (3'-cap) or may be present on both termini. In non-limiting examples: the 5'-cap is selected from the group comprising inverted abasic residue (moiety); 4',5'-methylene nucleotide; 1-(beta-D-erythrofuransyl) nucleotide; 4'-thio nucleotide; carbocyclic nucleotide; 1,5-

- anhydrohexitol nucleotide; L-nucleotides; alpha-nucleotides; modified base nucleotide; phosphorodithioate linkage; *threo*-pentofuranosyl nucleotide; acyclic 3',4'-seco nucleotide; acyclic 3,4-dihydroxybutyl nucleotide; acyclic 3,5-dihydroxypentyl nucleotide, 3'-3'-inverted nucleotide moiety; 3'-3'-inverted abasic moiety; 3'-2'-inverted nucleotide moiety; 3'-2'-inverted abasic moiety; 1,4-butanediol phosphate; 3'-phosphoramidate; hexylphosphate; aminohexyl phosphate; 3'-phosphate; 3'-phosphorothioate; phosphorodithioate; or bridging or non-bridging methylphosphonate moiety (for more details, see Wincott *et al.*, International PCT publication No. WO 97/26270, incorporated by reference herein).
- 10 In yet another preferred embodiment, the 3'-cap is selected from a group comprising, 4',5'-methylene nucleotide; 1-(beta-D-erythrofuransyl) nucleotide; 4'-thio nucleotide, carbocyclic nucleotide; 5'-amino-alkyl phosphate; 1,3-diamino-2-propyl phosphate; 3-aminopropyl phosphate; 6-aminohexyl phosphate; 1,2-aminododecyl phosphate; hydroxypropyl phosphate; 1,5-anhydrohexitol nucleotide; L-nucleotide; alpha-
- 15 nucleotide; modified base nucleotide; phosphorodithioate; *threo*-pentofuranosyl nucleotide; acyclic 3',4'-seco nucleotide; 3,4-dihydroxybutyl nucleotide; 3,5-dihydroxypentyl nucleotide, 5'-5'-inverted nucleotide moiety; 5'-5'-inverted abasic moiety; 5'-phosphoramidate; 5'-phosphorothioate; 1,4-butanediol phosphate; 5'-amino; bridging and/or non-bridging 5'-phosphoramidate, phosphorothioate and/or phosphorodithioate,
- 20 bridging or non bridging methylphosphonate and 5'-mercapto moieties (for more details see Beaucage and Iyer, 1993, *Tetrahedron* 49, 1925; incorporated by reference herein).

An "alkyl" group refers to a saturated aliphatic hydrocarbon, including straight-chain, branched-chain, and cyclic alkyl groups. Preferably, the alkyl group has 1 to 12 carbons. More preferably it is a lower alkyl of from 1 to 7 carbons, more preferably 1 to 4

25 carbons. The alkyl group may be substituted or unsubstituted. When substituted the substituted group(s) is preferably, hydroxyl, cyano, alkoxy, =O, =S, NO₂ or N(CH₃)₂, amino, or SH. The term also includes alkenyl groups which are unsaturated hydrocarbon groups containing at least one carbon-carbon double bond, including straight-chain, branched-chain, and cyclic groups. Preferably, the alkenyl group has 1 to 12 carbons.

30 More preferably it is a lower alkenyl of from 1 to 7 carbons, more preferably 1 to 4 carbons. The alkenyl group may be substituted or unsubstituted. When substituted the

substituted group(s) is preferably, hydroxyl, cyano, alkoxy, =O, =S, NO₂, halogen, N(CH₃)₂, amino, or SH. The term "alkyl" also includes alkynyl groups which have an unsaturated hydrocarbon group containing at least one carbon-carbon triple bond, including straight-chain, branched-chain, and cyclic groups. Preferably, the alkynyl group has 1 to 12 carbons. More preferably it is a lower alkynyl of from 1 to 7 carbons, more preferably 1 to 4 carbons. The alkynyl group may be substituted or unsubstituted. When substituted the substituted group(s) is preferably, hydroxyl, cyano, alkoxy, =O, =S, NO₂ or N(CH₃)₂, amino or SH.

Such alkyl groups may also include aryl, alkylaryl, carbocyclic aryl, heterocyclic aryl, amide and ester groups. An "aryl" group refers to an aromatic group which has at least one ring having a conjugated pi electron system and includes carbocyclic aryl, heterocyclic aryl and biaryl groups, all of which may be optionally substituted. The preferred substituent(s) of aryl groups are halogen, trihalomethyl, hydroxyl, SH, OH, cyano, alkoxy, alkyl, alkenyl, alkynyl, and amino groups. An "alkylaryl" group refers to an alkyl group (as described above) covalently joined to an aryl group (as described above). Carbocyclic aryl groups are groups wherein the ring atoms on the aromatic ring are all carbon atoms. The carbon atoms are optionally substituted. Heterocyclic aryl groups are groups having from 1 to 3 heteroatoms as ring atoms in the aromatic ring and the remainder of the ring atoms are carbon atoms. Suitable heteroatoms include oxygen, sulfur, and nitrogen, and include furanyl, thienyl, pyridyl, pyrrolyl, N-lower alkyl pyrrolo, pyrimidyl, pyrazinyl, imidazolyl and the like, all optionally substituted. An "amide" refers to an -C(O)-NH-R, where R is either alkyl, aryl, alkylaryl or hydrogen. An "ester" refers to an -C(O)-OR', where R is either alkyl, aryl, alkylaryl or hydrogen.

By "nucleotide" as used herein is as recognized in the art to include natural bases (standard), and modified bases well known in the art. Such bases are generally located at the 1' position of a nucleotide sugar moiety. Nucleotides generally comprise a base, sugar and a phosphate group. The nucleotides can be unmodified or modified at the sugar, phosphate and/or base moiety, (also referred to interchangeably as nucleotide analogs, modified nucleotides, non-natural nucleotides, non-standard nucleotides and other; see, for example, Usman and McSwiggen, *supra*; Eckstein *et al.*, International PCT Publication No. WO 92/07065; Usman *et al.*, International PCT Publication No. WO 93/15187;

Uhlman & Peyman, *supra*, all are hereby incorporated by reference herein). There are several examples of modified nucleic acid bases known in the art as summarized by Limbach *et al.*, 1994, *Nucleic Acids Res.* 22, 2183. Some of the non-limiting examples of base modifications that can be introduced into nucleic acid molecules include, inosine,
5 purine, pyridin-4-one, pyridin-2-one, phenyl, pseudouracil, 2, 4, 6-trimethoxy benzene, 3-methyl uracil, dihydrouridine, naphthyl, aminophenyl, 5-alkylcytidines (*e.g.*, 5-methylcytidine), 5-alkyluridines (*e.g.*, ribothymidine), 5-halouridine (*e.g.*, 5-bromouridine) or 6-azapyrimidines or 6-alkylpyrimidines (*e.g.* 6-methyluridine), propyne, and others (Burgin *et al.*, 1996, *Biochemistry*, 35, 14090; Uhlman & Peyman,
10 *supra*).

By "modified bases" in this aspect is meant nucleotide bases other than adenine, guanine, cytosine and uracil at 1' position or their equivalents; such bases may be used at any position, for example, within the catalytic core of an enzymatic nucleic acid molecule and/or in the substrate-binding regions of the nucleic acid molecule. Such modified
15 nucleotides include dideoxynucleotides which have pharmaceutical utility well known in the art, as well as utility in basic molecular biology methods such as sequencing.

In a preferred embodiment, the invention features modified ribozymes with phosphate backbone modifications comprising one or more phosphorothioate, phosphorodithioate, methylphosphonate, morpholino, amidate carbamate, carboxymethyl,
20 acetamidate, polyamide, sulfonate, sulfonamide, sulfamate, formacetal, thioformacetal, and/or alkylsilyl, substitutions. For a review of oligonucleotide backbone modifications, see Hunziker and Leumann, 1995, *Nucleic Acid Analogues: Synthesis and Properties*, in *Modern Synthetic Methods*, VCH, 331-417, and Mesmaeker *et al.*, 1994, *Novel Backbone Replacements for Oligonucleotides*, in *Carbohydrate Modifications in Antisense Research*,
25 ACS, 24-39. These references are hereby incorporated by reference herein.

By "abasic" is meant sugar moieties lacking a base or having other chemical groups in place of a base at the 1' position, (for more details, see Wincott *et al.*, International PCT publication No. WO 97/26270).

By "unmodified nucleoside" or "unmodified nucleotide" is meant one of the bases
30 adenine, cytosine, guanine, thymine, uracil joined to the 1' carbon of β -D-ribo-furanose.

By "modified nucleoside" or "modified nucleotide" is meant any nucleotide base which contains a modification in the chemical structure of an unmodified nucleotide base, sugar and/or phosphate.

5 In connection with 2'-modified nucleotides as described for the present invention, by "amino" is meant 2'-NH₂ or 2'-O- NH₂, which may be modified or unmodified. Such modified groups are described, for example, in Eckstein et al., U.S. Patent 5,672,695 and Matulic-Adamic et al., WO 98/28317, which are both incorporated by reference in their entireties.

10 Various modifications to nucleic acid (e.g., antisense and ribozyme) structure can be made to enhance the utility of these molecules. Such modifications will enhance shelf-life, half-life *in vitro*, stability, and ease of introduction of such oligonucleotides to the target site, e.g., to enhance penetration of cellular membranes, and confer the ability to recognize and bind to targeted cells.

15 Use of these molecules will lead to better treatment of the disease progression by affording the possibility of combination therapies (e.g., multiple ribozymes targeted to different genes, ribozymes coupled with known small molecule inhibitors, or intermittent treatment with combinations of ribozymes (including different ribozyme motifs) and/or other chemical or biological molecules). The treatment of patients with nucleic acid molecules may also include combinations of different types of nucleic acid molecules.
20 Therapies may be devised which include a mixture of ribozymes (including different ribozyme motifs), antisense and/or 2-5A chimera molecules to one or more targets to alleviate symptoms of a disease.

Administration of Nucleic Acid Molecules

25 Methods for the delivery of nucleic acid molecules are described in Akhtar *et al.*, 1992, *Trends Cell Bio.*, 2, 139; and *Delivery Strategies for Antisense Oligonucleotide Therapeutics*, ed. Akhtar, 1995 which are both incorporated herein by reference. Sullivan *et al.*, PCT WO 94/02595, further describes the general methods for delivery of enzymatic RNA molecules. These protocols may be utilized for the delivery of virtually any nucleic
30 acid molecule. Nucleic acid molecules may be administered to cells by a variety of methods known to those familiar to the art, including, but not restricted to, encapsulation in liposomes, by iontophoresis, or by incorporation into other vehicles, such as hydrogels,

cyclodextrins, biodegradable nanocapsules, and bioadhesive microspheres. For some indications, nucleic acid molecules may be directly delivered *ex vivo* to cells or tissues with or without the aforementioned vehicles. Alternatively, the nucleic acid/vehicle combination is locally delivered by direct injection or by use of a catheter, infusion pump or stent. Many examples in the art describe CNS delivery methods of oligonucleotides by osmotic pump, (see Chun *et al.*, 1998, *Neuroscience Letters*, 257, 135-138, D'Aldin *et al.*, 1998, *Mol. Brain Research*, 55, 151-164, Dryden *et al.*, 1998, *J. Endocrinol.*, 157, 169-175, Ghirmikar *et al.*, 1998, *Neuroscience Letters*, 247, 21-24) or direct infusion (Broadus *et al.*, 1997, *Neurosurg. Focus*, 3, article 4). Other routes of delivery include, but are not limited to oral (tablet or pill form) and/or intrathecal delivery (Gold, 1997, *Neuroscience*, 76, 1153-1158). For a comprehensive review on drug delivery strategies including broad coverage of CNS delivery, see Jain, *Drug Delivery Systems: Technologies and Commercial Opportunities*, Decision Resources, 1998. Other routes of delivery include, but are not limited to, intravascular, intramuscular, subcutaneous or joint injection, aerosol inhalation, oral (tablet or pill form), topical, systemic, ocular, intraperitoneal and/or intrathecal delivery. More detailed descriptions of nucleic acid delivery and administration are provided in Sullivan *et al.*, *supra*, Draper *et al.*, PCT WO93/23569; Beigelman *et al.*, PCT WO99/05094, and Klimuk *et al.*, PCT WO99/04819 all of which are incorporated by reference herein.

The molecules of the instant invention can be used as pharmaceutical agents. Pharmaceutical agents prevent, inhibit the occurrence, or treat (alleviate a symptom to some extent, preferably all of the symptoms) of a disease state in a patient.

The negatively charged polynucleotides of the invention can be administered (*e.g.*, RNA, DNA or protein) and introduced into a patient by any standard means, with or without stabilizers, buffers, and the like, to form a pharmaceutical composition. When it is desired to use a liposome delivery mechanism, standard protocols for formation of liposomes can be followed. The compositions of the present invention may also be formulated and used as tablets, capsules or elixirs for oral administration; suppositories for rectal administration; sterile solutions; suspensions for injectable administration; and the other compositions known in the art.

The present invention also includes pharmaceutically acceptable formulations of the compounds described. These formulations include salts of the above compounds, *e.g.*, acid addition salts, for example, salts of hydrochloric, hydrobromic, acetic acid, and benzene sulfonic acid.

5 A pharmacological composition or formulation refers to a composition or formulation in a form suitable for administration, *e.g.*, systemic administration, into a cell or patient, preferably a human. Suitable forms, in part, depend upon the use or the route of entry, for example, oral, transdermal, or by injection. Such forms should not prevent the composition or formulation from reaching a target cell (*i.e.*, a cell to which the negatively
10 charged polymer is desired to be delivered to). For example, pharmacological compositions injected into the blood stream should be soluble. Other factors are known in the art, and include considerations such as toxicity and forms which prevent the composition or formulation from exerting its effect.

By "systemic administration" is meant *in vivo* systemic absorption or accumulation
15 of drugs in the blood stream followed by distribution throughout the entire body. Administration routes which lead to systemic absorption include, without limitations: intravenous, subcutaneous, intraperitoneal, inhalation, oral, intrapulmonary and intramuscular. Each of these administration routes expose the desired negatively charged polymers, *e.g.*, nucleic acids, to an accessible diseased tissue. The rate of entry of a drug
20 into the circulation has been shown to be a function of molecular weight or size. The use of a liposome or other drug carrier comprising the compounds of the instant invention can potentially localize the drug, for example, in certain tissue types, such as the tissues of the reticular endothelial system (RES). A liposome formulation which can facilitate the association of drug with the surface of cells, such as, lymphocytes and macrophages is also
25 useful. This approach may provide enhanced delivery of the drug to target cells by taking advantage of the specificity of macrophage and lymphocyte immune recognition of abnormal cells, such as cancer cells.

By pharmaceutically acceptable formulation is meant, a composition or formulation that allows for the effective distribution of the nucleic acid molecules of the instant
30 invention in the physical location most suitable for their desired activity. Nonlimiting examples of agents suitable for formulation with the nucleic acid molecules of the instant invention include: P-glycoprotein inhibitors (such as Pluronic P85) which can enhance

entry of drugs into the CNS (Jolliet-Riant and Tillement, 1999, *Fundam. Clin. Pharmacol.*, 13, 16-26); biodegradable polymers, such as poly (DL-lactide-coglycolide) microspheres for sustained release delivery after intracerebral implantation (Emerich, DF *et al.*, 1999, *Cell Transplant*, 8, 47-58) Alkermes, Inc. Cambridge, MA; and loaded nanoparticles, such as those made of polybutylcyanoacrylate, which can deliver drugs across the blood brain barrier and can alter neuronal uptake mechanisms (*Prog Neuropsychopharmacol Biol Psychiatry*, 23, 941-949, 1999). Other non-limiting examples of delivery strategies for the nucleic acid molecules of the instant invention include material described in Boado *et al.*, 1998, *J. Pharm. Sci.*, 87, 1308-1315; Tyler *et al.*, 1999, *FEBS Lett.*, 421, 280-284; Pardridge *et al.*, 1995, *PNAS USA.*, 92, 5592-5596; Boado, 1995, *Adv. Drug Delivery Rev.*, 15, 73-107; Aldrian-Herrada *et al.*, 1998, *Nucleic Acids Res.*, 26, 4910-4916; and Tyler *et al.*, 1999, *PNAS USA.*, 96, 7053-7058.

The invention also features the use of the composition comprising surface-modified liposomes containing poly (ethylene glycol) lipids (PEG-modified, or long-circulating liposomes or stealth liposomes). These formulations offer a method for increasing the accumulation of drugs in target tissues. This class of drug carriers resists opsonization and elimination by the mononuclear phagocytic system (MPS or RES), thereby enabling longer blood circulation times and enhanced tissue exposure for the encapsulated drug (Lasic *et al. Chem. Rev.* 1995, 95, 2601-2627; Ishiwata *et al.*, *Chem. Pharm. Bull.* 1995, 43, 1005-1011). Such liposomes have been shown to accumulate selectively in tumors, presumably by extravasation and capture in the neovascularized target tissues (Lasic *et al.*, *Science* 1995, 267, 1275-1276; Oku *et al.*, 1995, *Biochim. Biophys. Acta*, 1238, 86-90). The long-circulating liposomes enhance the pharmacokinetics and pharmacodynamics of DNA and RNA, particularly compared to conventional cationic liposomes which are known to accumulate in tissues of the MPS (Liu *et al.*, *J. Biol. Chem.* 1995, 42, 24864-24870; Choi *et al.*, International PCT Publication No. WO 96/10391; Ansell *et al.*, International PCT Publication No. WO 96/10390; Holland *et al.*, International PCT Publication No. WO 96/10392; all of which are incorporated herein by reference). Long-circulating liposomes are also likely to protect drugs from nuclease degradation to a greater extent compared to cationic liposomes, based on their ability to avoid accumulation in metabolically aggressive MPS tissues such as the liver and spleen.

The present invention also includes compositions prepared for storage or administration which include a pharmaceutically effective amount of the desired compounds in a pharmaceutically acceptable carrier or diluent. Acceptable carriers or diluents for therapeutic use are well known in the pharmaceutical art, and are described, for example, in *Remington's Pharmaceutical Sciences*, Mack Publishing Co. (A.R. Gennaro edit. 1985) hereby incorporated by reference herein. For example, preservatives, stabilizers, dyes and flavoring agents may be provided. These include sodium benzoate, sorbic acid and esters of *p*-hydroxybenzoic acid. In addition, antioxidants and suspending agents may be used.

A pharmaceutically effective dose is that dose required to prevent, inhibit the occurrence, or treat (alleviate a symptom to some extent, preferably all of the symptoms) of a disease state. The pharmaceutically effective dose depends on the type of disease, the composition used, the route of administration, the type of mammal being treated, the physical characteristics of the specific mammal under consideration, concurrent medication, and other factors which those skilled in the medical arts will recognize. Generally, an amount between 0.1 mg/kg and 100 mg/kg body weight/day of active ingredients is administered dependent upon potency of the negatively charged polymer.

The nucleic acid molecules of the present invention may also be administered to a patient in combination with other therapeutic compounds to increase the overall therapeutic effect. The use of multiple compounds to treat an indication may increase the beneficial effects while reducing the presence of side effects.

Alternatively, certain of the nucleic acid molecules of the instant invention can be expressed within cells from eukaryotic promoters (*e.g.*, Izant and Weintraub, 1985, *Science*, 229, 345; McGarry and Lindquist, 1986, *Proc. Natl. Acad. Sci.*, USA 83, 399; Scanlon *et al.*, 1991, *Proc. Natl. Acad. Sci. USA*, 88, 10591-5; Kashani-Sabet *et al.*, 1992, *Antisense Res. Dev.*, 2, 3-15; Dropulic *et al.*, 1992, *J. Virol.*, 66, 1432-41; Weerasinghe *et al.*, 1991, *J. Virol.*, 65, 5531-4; Ojwang *et al.*, 1992, *Proc. Natl. Acad. Sci. USA*, 89, 10802-6; Chen *et al.*, 1992, *Nucleic Acids Res.*, 20, 4581-9; Sarver *et al.*, 1990 *Science*, 247, 1222-1225; Thompson *et al.*, 1995, *Nucleic Acids Res.*, 23, 2259; Good *et al.*, 1997, *Gene Therapy*, 4, 45; all of these references are hereby incorporated herein, in their totalities, by reference). Those skilled in the art realize that any nucleic acid can be expressed in eukaryotic cells from the appropriate DNA/RNA vector. The activity of such

nucleic acids can be augmented by their release from the primary transcript by a ribozyme (Draper *et al.*, PCT WO 93/23569, and Sullivan *et al.*, PCT WO 94/02595; Ohkawa *et al.*, 1992, *Nucleic Acids Symp. Ser.*, 27, 15-6; Taira *et al.*, 1991, *Nucleic Acids Res.*, 19, 5125-30; Ventura *et al.*, 1993, *Nucleic Acids Res.*, 21, 3249-55; Chowrira *et al.*, 1994, *J. Biol. Chem.*, 269, 25856; all of these references are hereby incorporated in their totality by reference herein).

In another aspect of the invention, RNA molecules of the present invention are preferably expressed from transcription units (see, for example, Couture *et al.*, 1996, *TIG.*, 12, 510) inserted into DNA or RNA vectors. The recombinant vectors are preferably DNA plasmids or viral vectors. Ribozyme expressing viral vectors could be constructed based on, but not limited to, adeno-associated virus, retrovirus, adenovirus, or alphavirus. Preferably, the recombinant vectors capable of expressing the nucleic acid molecules are delivered as described above, and persist in target cells. Alternatively, viral vectors may be used that provide for transient expression of nucleic acid molecules. Such vectors might be repeatedly administered as necessary. Once expressed, the nucleic acid molecule binds to the target mRNA. Delivery of nucleic acid molecule expressing vectors could be systemic, such as by intravenous or intra-muscular administration, by administration to target cells ex-planted from the patient followed by reintroduction into the patient, or by any other means that would allow for introduction into the desired target cell (for a review see Couture *et al.*, 1996, *TIG.*, 12, 510).

In one aspect, the invention features an expression vector comprising a nucleic acid sequence encoding at least one of the nucleic acid molecules of the instant invention is disclosed. The nucleic acid sequence encoding the nucleic acid molecule of the instant invention is operably linked in a manner which allows expression of that nucleic acid molecule.

In another aspect the invention features an expression vector comprising: a) a transcription initiation region (*e.g.*, eukaryotic pol I, II or III initiation region); b) a transcription termination region (*e.g.*, eukaryotic pol I, II or III termination region); c) a nucleic acid sequence encoding at least one of the nucleic acid catalyst of the instant invention; and wherein said sequence is operably linked to said initiation region and said termination region, in a manner which allows expression and/or delivery of said nucleic acid molecule. The vector may optionally include an open reading frame (ORF) for a

protein operably linked on the 5' side or the 3'-side of the sequence encoding the nucleic acid catalyst of the invention; and/or an intron (intervening sequences).

Transcription of the nucleic acid molecule sequences are driven from a promoter for eukaryotic RNA polymerase I (pol I), RNA polymerase II (pol II), or RNA polymerase III (pol III). Transcripts from pol II or pol III promoters will be expressed at high levels in all
5 cells; the levels of a given pol II promoter in a given cell type will depend on the nature of the gene regulatory sequences (enhancers, silencers, etc.) present nearby. Prokaryotic RNA polymerase promoters are also used, providing that the prokaryotic RNA polymerase enzyme is expressed in the appropriate cells (Elroy-Stein and Moss, 1990, *Proc. Natl. Acad. Sci. U S A*, 87, 6743-7; Gao and Huang 1993, *Nucleic Acids Res.*, 21, 2867-72; Lieber et al., 1993, *Methods Enzymol.*, 217, 47-66; Zhou et al., 1990, *Mol. Cell. Biol.*, 10, 4529-37). All of these references are incorporated by reference herein. Several investigators have demonstrated that nucleic acid molecules, such as ribozymes expressed from such promoters can function in mammalian cells (e.g. Kashani-Sabet et al., 1992, *Antisense Res. Dev.*, 2, 3-15; Ojwang et al., 1992, *Proc. Natl. Acad. Sci. U S A*, 89, 10802-6; Chen et al., 1992, *Nucleic Acids Res.*, 20, 4581-9; Yu et al., 1993, *Proc. Natl. Acad. Sci. U S A*, 90, 6340-4; L'Huillier et al., 1992, *EMBO J.*, 11, 4411-8; Lisiewicz et al., 1993, *Proc. Natl. Acad. Sci. U. S. A*, 90, 8000-4; Thompson et al., 1995, *Nucleic Acids Res.*, 23, 2259; Sullenger & Cech, 1993, *Science*, 262, 1566). More specifically,
15 transcription units such as the ones derived from genes encoding U6 small nuclear (snRNA), transfer RNA (tRNA) and adenovirus VA RNA are useful in generating high concentrations of desired RNA molecules such as ribozymes in cells (Thompson et al., *supra*; Couture and Stinchcomb, 1996, *supra*; Noonberg et al., 1994, *Nucleic Acid Res.*, 22, 2830; Noonberg et al., US Patent No. 5,624,803; Good et al., 1997, *Gene Ther.*, 4, 45; Beigelman et al., International PCT Publication No. *WO 96/18736*; all of these
25 publications are incorporated by reference herein. The above ribozyme transcription units can be incorporated into a variety of vectors for introduction into mammalian cells, including but not restricted to, plasmid DNA vectors, viral DNA vectors (such as adenovirus or adeno-associated virus vectors), or viral RNA vectors (such as retroviral or
30 alphavirus vectors) (for a review see Couture and Stinchcomb, 1996, *supra*).

In yet another aspect, the invention features an expression vector comprising nucleic acid sequence encoding at least one of the nucleic acid molecules of the invention, in a manner which allows expression of that nucleic acid molecule. The expression vector comprises in one embodiment; a) a transcription initiation region; b) a transcription termination region; c) a nucleic acid sequence encoding at least one said nucleic acid molecule; and wherein said sequence is operably linked to said initiation region and said termination region, in a manner which allows expression and/or delivery of said nucleic acid molecule. In another preferred embodiment the expression vector comprises: a) a transcription initiation region; b) a transcription termination region; c) an open reading frame; d) a nucleic acid sequence encoding at least one said nucleic acid molecule, wherein said sequence is operably linked to the 3'-end of said open reading frame; and wherein said sequence is operably linked to said initiation region, said open reading frame and said termination region, in a manner which allows expression and/or delivery of said nucleic acid molecule. In yet another embodiment, the expression vector comprises: a) a transcription initiation region; b) a transcription termination region; c) an intron; d) a nucleic acid sequence encoding at least one said nucleic acid molecule; and wherein said sequence is operably linked to said initiation region, said intron and said termination region, in a manner which allows expression and/or delivery of said nucleic acid molecule. In another embodiment, the expression vector comprises: a) a transcription initiation region; b) a transcription termination region; c) an intron; d) an open reading frame; e) a nucleic acid sequence encoding at least one said nucleic acid molecule, wherein said sequence is operably linked to the 3'-end of said open reading frame; and wherein said sequence is operably linked to said initiation region, said intron, said open reading frame and said termination region, in a manner which allows expression and/or delivery of said nucleic acid molecule.

Examples:

The following are non-limiting examples showing the selection, isolation, synthesis and activity of nucleic acids of the instant invention.

Example 1: Telomerase

The ribonucleoprotein enzyme telomerase consists of an RNA template subunit and one or more protein subunits including telomerase reverse transcriptase (TERT), which function together to direct the synthesis of telomeres. Telomeres exist as non-nucleosome DNA/protein complexes at the physical ends of eukaryotic chromosomes. These capping structures maintain chromosome stability and replicative potential (Zakian, V. A., 1995, Science, 270, 1601-1607). Telomere structure is characterized by tandem repeats of conserved DNA sequences rich in G-C base pairs. Additional conserved telomere elements include a terminal 3'-overhang in the G-rich strand and non-histone structural proteins that are complexed with telomeric DNA in the nucleus. (Blackburn, "E., 1990, JBC., 265, 5919-5921.). Observed shortening of telomeres coincides with the onset of cellular senescence in most somatic cell lines lacking significant levels of telomerase. This finding has had a profound impact on our views concerning the mechanisms of aging, age related disease, and cancer.

Conventional DNA polymerases are unable to fully replicate the ends of linear chromosomes (Watson, J. D., 1972, Nature, 239, 197-201). This inability stems from the 3' G-rich overhang that is a product of ribonuclease cleavage of the RNA primer used in DNA replication. The overhang prevents DNA polymerase replication since the recessed C-rich parent strand cannot be used as a template. Telomerase overcomes this limitation by extending the 3' end of the chromosome using deoxyribonucleotides as substrates and a sequence within the telomerase RNA subunit as a template. (Lingner, J., 1995, Science, 269, 1533-1534). As such, telomerase is considered a reverse transcriptase that is responsible for telomere maintenance.

Telomerase was first discovered by in *Tetrahymena thermophila* in 1985 (Greider, C. W., 1995, Cell, 43, 405-413). The RNA subunits and their respective genes were later discovered and characterized in protozoa, budding yeast, and mammals. Genetic studies of these genes confirmed the role of telomerase RNA (TR) in determining telomere sequence by mutating genes which encode the telomeric RNA (Yu, G. L., 1990, Nature, 344, 126-132), (Singer, M. S., 1994, Science, 266, 404-409), (Blasco, M. A., 1995, Science, 269, 1267-1270). These studies showed that telomerase activity parallels TR expression in protozoa, yeast and mice. However, the expression of human telomerase RNA (hTR) does not correlate well with telomerase activity in mammalian cells. Many

human tissues express hTR but are devoid of telomerase activity (Feng, J., 1995, Science, 269, 1236-1241). Knockout mice, in which the mTR gene has been deleted from germline cells, have been shown to be viable for at least six generations. Cells from later generations of these mice showed chromosomal abnormalities consistent with telomere degradation, indicating that mTR is necessary for telomere length maintenance, but is not required for embryonic development, oncogenic transformation, or tumor formation in mice (Blasco, M. A., 1997, Cell, 91, 25-34).

The first catalytically active subunit of telomerase (p123) was isolated from *Euplotes aediculatus* along with another subunit (p43) and a 66-kD RNA subunit (Linger, J., 1996, Proc. Natl. Acad. Sci., 93, 10712-10717). Subsequent studies revealed telomerase catalytic subunit homologs from fission yeast (Est2p) and human genes (TRT1). The human homolog, TRT1 encoding hTERT, expressed mRNA with a strong correlation to telomerase activity in human cells (Nakamura, T. M., 1997, Science, 277, 955-959). Reconstitution of telomerase activity with *in vitro* transcribed and translated hTERT and hTR, either co-synthesized or simply mixed, demonstrated that hTERT and hTR represent the minimal components of telomerase. Furthermore, transient expression of hTERT in normal diploid human cells restored telomerase activity, demonstrating that hTERT is the only component necessary to restore telomerase activity in normal human cells (Weinrich, S. L., 1997, Nature Genetics, 17, 498-502). The introduction of telomerase into normal human cells using hTERT expression via transfection has resulted in the extension of life span in these cells. Such findings indicate that telomere loss in the absence of telomerase is the "mitotic clock" that controls the replicative potential of a cell prior to senescence (Bodnar, A. G., 1998, Science, 279, 349-352).

Expression of telomerase is observed in germ cell and most cancer cell lines. These "immortal" cell lines continue to divide without shortening of their telomeres (Kim, N. W., 1994, Science, 266, 2011-2015). A model of tumor progression has evolved from these findings, suggesting a role for telomerase expression in malignant transformation. Successful malignant transformation in human cells was accomplished for the first time by ectopic expression of hTERT in combination with two oncogenes, SV40 large-T and H-ras. Injection of nude mice with cells expressing these oncogenes and hTERT resulted in rapid growth of tumors. These observations indicate that hTERT mediated telomere

maintenance is essential for the formation of human tumor cells (Hahn, W. C., 1999, Nature, 400, 464-468).

Various methods have been developed to assay telomerase activity *in vitro*. The most widely used method to characterize telomerase activity is the telomeric repeat amplification protocol (TRAP). TRAP utilizes RT-PCR of cellular extracts to measure telomerase activity by making the amount of PCR target dependant upon the biochemical activity of the enzyme (Kim, N. W., 1997, Nucleic Acids Research, 25, 2595-2597, which is incorporated by reference herein).

A method based on Kim is as follows. Briefly, for the telomerase assay, 2µg of protein extract is used. The extract is assayed in 50µl of reaction mixture containing 0.1 µg TS substrate primer (5'-AATCCGTCGAGCAGAGTT-3', end-labeled using alpha-³²P-ATP and T4 polynucleotide kinase), 0.1µg ACX return primer(5'-GCGCGG[CTTACC]₃CTAACC-3'), 0.1 µg NT internal control primer (5'-ATCGCTTCTCGGCCTTTT-3'), 0.01 micromol TSNT internal control template (5'-AATCCGTCGAGCAGAGTTAAAAGGCCGAGAACGAT-3'), 50 µM each deoxynucleoside triphosphate, 2 U of Taq DNA polymerase, and 2 µl CHAPS protein extract, all in 1X TRAP buffer (20 mM Tris (pH 8.3), 68 mM KCl, 1.5 mM MgCl₂, 1 mM EGTA, 0.05% Tween 20). Each reaction is placed in a thermocycler block preheated to 30 C and incubated at 30 C for 10 minutes, then cycled for 27 cycles of 94 degrees C for 30 seconds, 60 degrees C for 30 seconds. Reaction products are separated on a denaturing 8% polyacrylamide gel, followed by drying of the gel and autoradiography. The internal control (to control for possible Taq polymerase inhibition) generates a band of 36 nt. Comparison of radioactive signal integrated (*e.g.*, by phosphorimager analysis) for telomerase-extended bands with the radioactive signal from a reaction performed with a known amount of quantification standard template (termed R8; 5'-AATCCGTCGAGCAGAGTTAG [GGTTAG]₇-3') allows expression of telomerase activity as an absolute value. The absolute value = $\frac{\text{TPG (total product generated)}}{[(\text{TP} - \text{TPi})/\text{TI}]/[(\text{R8-B})/\text{RI}]} \times 100$, where TP = telomerase products from test extract, TPi = telomerase products from a heat-inactivated (75 C, 10 minutes) extract reaction, TI = the signal from the internal control, R8 = the signal from the R8 qualification standard template reaction, B = signal from a lysis buffer-only blank reaction, and RI = the internal control value for the reaction containing R8 template and NT and TSNT control primers.

TPG values of 0-10,000 are possible, with the linear range being from approximately 1 to 1000 TPG. The range of 1 to 1000 TPG encompasses the minimum and maximum levels of telomerase activity in most tumor samples tested, while non-tumor cells most often have no telomerase activity (TPG approximately zero).

- 5 Telomerase activity may also be assayed as follows. Samples to be assayed for telomerase activity are prepared by extraction into CHAPS lysis buffer (10mM Tris pH 7.5, 1mM MgCl₂, 1mM EGTA, 0.1 mM PMSF, 5mM -mercaptoethanol, 1mM DTT, 0.5% 3-[(3-cholamidopropyl)-dimethyl-amino]-1- propanesulfonate (CHAPS), 10% glycerol and 40 U/ml RNase inhibitor (Promega, Madison, WI, U.S.A.). Cells are
- 10 suspended in CHAPS lysis buffer and incubated on ice for 30 minutes, which allows lysis of 90-100% of cells. Lysate is then transferred to polyallomer centrifuge tubes and spun at 100,000 x g for 1 hour at 4 degrees C. The supernatant is the protein extract, and concentration ranges of 4-10 µg/µl are suitable for telomerase assay. Extracts may be concentrated if necessary using a Microcon Microfilter 30 (Amicon, Beverly, MA U.S.A.)
- 15 according to the manufacturer's instructions. Extracts may be stored frozen at -80 degrees C until assayed.

- A variety of animal models have been designed to assay telomerase activity *in vivo*. Inhibition of telomerase activity has been analyzed in rats via cell proliferation studies with MNU (N-methyl-N-nitrosurea) induced mammary carcinomas in response to treatment
- 20 with 4-(hydroxyphenyl)retinamide (4-HPR), a known inhibitor of mammary carcinogenesis in animal models and premenopausal women (Bednarek, A., 1999, Carcinogenesis, 20, 879-883). Additional studies have focused on the up-regulation of telomerase in transformed cell lines from animal and human model systems (Zhang, P. B., 1998, Leuk. Res., 22, 509-516), (Chadeneau, C., 1995, Oncogene, 11, 893-898),
- 25 (Greenberg, R., 1999, Oncogene, 18, 1219-1226).

- Human cell culture studies have been established to assay inhibition of telomerase activity in human carcinomas responding to various therapeutics. A human breast cancer model for studying telomerase inhibitors is described (Raymond, E., 1999, Br. J. Cancer, 80, 1332-1341). Human studies of telomerase expression as related to various other
- 30 cancers are described including cervical cancer (Nakano, K., 1998, Am. J. Pathol, 153, 857-864), endometrial cancer (Kyo, S., 1999, Int. J. Cancer, 80, 60-63), meningeal carcinoma (Kleinschmidt-DeMasters, B. K., 1998, J. Neurol. Sci., 161, 124-134), lung

carcinoma (Yashima, K., 1997, Cancer Reseach, 57, 2372-2377), testicular cancer in response to cisplatin (Burger, A. M., 1997, Eur. J. Cancer, 33, 638-644), and ovarian carcinoma (Counter, C. M., 1994, Proc. Natl. Acad. Sci., 91, 2900-2904).

Particular degenerative and disease states that can be associated with telomerase expression modulation include but are not limited to:

Cancer: Almost all human tumors have detectable telomerase activity (Shay, J. W., 1997, Eur. J. Cancer, 33, 787-791). Treatment with telomerase inhibitors may provide effective cancer therapy with minimal side effects in normal somatic cells that lack telomerase activity. The therapeutic potential exists for the treatment of a wide variety of cancer types.

Restinosis: Telomerase inhibition in vascular smooth muscle cells may inhibit restinosis by limiting proliferation of these cells.

Infectious disease: Telomerase inhibition in infectious cell types that express telomerase activity may provide selective anti-infectious agent activity. Such treatment may prove especially effective in protozoan-based infection such as Giardia and Lesh Meniesis.

Transplant rejection: Telomerase inhibition in endothelial cell types may demonstrate selective immunnosuppressant activity. Activation of telomerase in transplant cells could benefit grafting success through increased proliferative potential.

Autoimmune disease: Telomerase modulation in various immune cells may prove beneficial in treating diseases such as multiple sclerosis, lupus, and AIDS.

Age related disease: Activation of telomerase expression in cells at or nearing senescence as a result of advanced age or premature aging could benefit conditions such as macular degeneration, skin ulceration, and rheumatoid arthritis.

The present body of knowledge in telomerase research indicates the need for methods to assay telomerase activity and for compounds that can regulate telomerase expression for research, diagnostic, trait alteration, animal health and therapeutic use.

Gemcytabine and cyclophosphamide are non-limiting examples of chemotherapeutic agents that can be combined with or used in conjunction with the nucleic acid molecules (e.g. ribozymes and antisense molecules) of the instant invention. Those skilled in the art will recognize that other drugs such as anti-cancer compounds and therapies can be similarly be readily combined with the nucleic acid molecules of the instant invention

(e.g. ribozymes and antisense molecules) and are hence within the scope of the instant invention. Such compounds and therapies are well known in the art (see for example *Cancer: Principles and Practice of Oncology*, Volumes 1 and 2, eds Devita, V.T., Hellman, S., and Rosenberg, S.A., J.B. Lippincott Company, Philadelphia, USA; incorporated herein by reference) and include, without limitations, antifolates; fluoropyrimidines; cytarabine; purine analogs; adenosine analogs; amsacrine; topoisomerase I inhibitors; anthracyclins; retinoids; antibiotics such as bleomycin, anthacyclins, mitomycin C, dactinomycin, and mithramycin; hexamethylmelamine; dacarbazine; l-asparaginase; platinum analogs; alkylating agents such as nitrogen mustard, melphalan, chlorambucil, busulfan, ifosfamide, 4-hydroperoxycyclophosphamide, nitrosoureas, thiotepa; plant derived compounds such as vinca alkaloids, epipodophyllotoxins, taxol; Tomaxifen; radiation therapy; surgery; nutritional supplements; gene therapy; radiotherapy such as 3D-CRT; immunotoxin therapy such as ricin, monoclonal antibodies herceptin; and the like. For combination therapy, the nucleic acids of the invention are prepared in one of two ways. First, the agents are physically combined in a preparation of nucleic acid and chemotherapeutic agent, such as a mixture of a nucleic acid of the invention encapsulated in liposomes and ifosfamide in a solution for intravenous administration, wherein both agents are present in a therapeutically effective concentration (e.g., ifosfamide in solution to deliver 1000-1250 mg/m²/day and liposome-associated nucleic acid of the invention in the same solution to deliver 0.1-100 mg/kg/day). Alternatively, the agents are administered separately but simultaneously in their respective effective doses (e.g., 1000-1250 mg/m²/d ifosfamide and 0.1 to 100 mg/kg/day nucleic acid of the invention).

Gaeta *et al.*, US patents No. 5,760,062; 5,767,278; 5,770,613 have described small molecule inhibitors of human telomerase RNA (hTR) subunit.

Blasco *et al.*, 1995, Science, 269, 1267-1270 describe the synthesis and testing of antisense oligonucleotides targeted against a specific region of the mouse telomerase RNA (mTR) subunit and reported reduction in telomerase activity in mice.

Bisoffi *et al.*, 1998, Eur. J. Cancer, 34, 1242-1249 have studied the down regulation of human telomerase activity by a retrovirus vector expressing antisense RNA targeted against the hTR RNA.

Norton *et al.*, 1996, Nature Biotechnology, 14, 615-619 have reported the use of a peptide nucleic acid (PNA) molecule targeting hTR RNA to down regulate telomerase activity in human immortal breast epithelial cells.

Yokoyama *et al.*, 1998, Cancer Research, 58, 5406-5410 have reported the
5 synthesis and testing of hammerhead ribozyme constructs targeting hTR RNA resulting in a decrease in the telomerase activity in Ishikawa cells.

Henderson, European Patent Application No. 666,313-A2 describes methods of identifying and cloning hTR gene for use in gene therapy approaches for creating aberrant telomeric sequences in transfected human tumor cells. A ribozyme based gene therapy
10 approach to inhibit the expression of hTR gene is described as well. The intended result of such therapies involves incurred genetic instability based on non-native telomeric sequences resulting in rapid cell death of the treated cells.

West *et al.*, US patent No. 5,489,508 describe methods for determining telomere length and telomerase activity in cells. Inhibitors of hTR RNA, including oligonucleotides
15 and/or small molecules are described.

These foregoing approaches of targeting the telomerase RNA subunit (TR) may not be very beneficial, because as demonstrated by Feng *et al.*, (Feng, J., 1995, Science, 269, 1236-1241), telomerase activity in humans does not correlate well to hTR concentration.

Collins *et al.*, International PCT publication No. WO 98/01542 describes assays for
20 the detection of telomerase activity. Four human telomerase subunit proteins are described called p140, p105, p48 and p43. In addition, hybridization probes and primers are described as inhibitors of telomerase gene function. Antibody based inhibitors of telomerase protein subunits are described.

A more attractive approach to telomerase regulation would involve the regulation of
25 human telomerase by modulating the expression of the protein subunits of the enzyme, preferably the reverse transcriptase (hTERT) subunit. Based on reconstitution experiments, hTERT and hTR represent the minimal components of telomerase. Since hTR expression does not correlate well with telomerase activity in human cells and since many human cells express hTR without telomerase activity, targeting hTERT may prove
30 more beneficial than targeting hTR. hTERT is the only component necessary to restore telomerase activity in normal human cells. A study in which the three major subunits of telomerase (hTR, TP1, and hTERT) were assayed in normal and malignant endometrial

tissues determined that hTERT is a rate limiting determinant of enzymatic activity of human telomerase (Kyo, S., 1999, Int. J. Cancer, 80, 60-63). Additional protein subunits that have been isolated most likely serve only a structural role in telomerase activity, but may be important in enhancing the activity of the telomerase enzyme. As such, hTERT is one of the better targets for the ectopic regulation of telomerase activity.

Cech *et al.*, International PCT publication No. WO 98/14593 describe compositions and methods related to hTERT for diagnosis, prognosis and treatment of human diseases, for altering proliferative capacity in cells and organisms, and for screening compounds and treatments with potential use as human therapeutics.

Cech *et al.*, International PCT publication No. WO 98/14592 describe nucleic acid and amino acid sequences encoding various telomerase protein subunits and motifs of *Euplotes aediculatus*, and related sequences from *Schizosaccharomyces*, *Saccharomyces* sequences, and human telomerase. The polypeptides comprising telomeric subunits and functional polypeptides and ribonucleoproteins that contain these subunits are described as well. Cech *et al.*, International PCT Publication No. WO 98/14592, mentions in general terms the possibility of using antisense and ribozymes to down regulate the expression of human telomerase reverse transcriptase enzyme.

Identification of Potential Target Sites in Human TERT RNA

The sequence of human TERT was screened for accessible sites using a computer folding algorithm. Regions of the RNA that did not form secondary folding structures and contained potential ribozyme and/or antisense binding/cleavage sites were identified. The sequences of these cleavage sites are shown in Tables 13-17.

Selection of Enzymatic Nucleic Acid Cleavage Sites in Human TERT RNA

To test whether the sites predicted by the computer-based RNA folding algorithm corresponded to accessible sites in TERT RNA, 10 hammerhead ribozyme and three G-Cleaver ribozyme sites were selected for further analysis (Table 17). Ribozyme target sites were chosen by analyzing sequences of Human TERT (Nakamura *et al.*, 1997 Science 277, 955-959; Genbank sequence accession number: NM_003219) and prioritizing the sites on the basis of folding. Ribozymes were designed that could bind each target and were individually analyzed by computer folding (Christoffersen *et al.*,

1994 *J. Mol. Struc. Theochem*, 311, 273; Jaeger *et al.*, 1989, *Proc. Natl. Acad. Sci. USA*, 86, 7706) to assess whether the ribozyme sequences fold into the appropriate secondary structure. Those ribozymes with unfavorable intramolecular interactions between the binding arms and the catalytic core were eliminated from consideration. As noted below, varying binding arm lengths can be chosen to optimize activity. Generally, at least 5 bases on each arm are able to bind to, or otherwise interact with, the target RNA.

Chemical Synthesis and Purification of Ribozymes for Efficient Cleavage of TERT RNA

Ribozymes were designed to anneal to various sites in the RNA message. The binding arms are complementary to the target site sequences described above. The ribozymes were chemically synthesized. The method of synthesis used followed the procedure for normal RNA synthesis as described above and in Usman *et al.*, (1987 *J. Am. Chem. Soc.*, 109, 7845), Scaringe *et al.*, (1990 *Nucleic Acids Res.*, 18, 5433) and Wincott *et al.*, *supra*, and made use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. The average stepwise coupling yields were >98%.

Ribozymes were also synthesized from DNA templates using bacteriophage T7 RNA polymerase (Milligan and Uhlenbeck, 1989, *Methods Enzymol.* 180, 51). Ribozymes were purified by gel electrophoresis using general methods or were purified by high pressure liquid chromatography (HPLC; See Wincott *et al.*, *supra*; the totality of which is hereby incorporated herein by reference) and were resuspended in water. The sequences of the chemically synthesized ribozymes used in this study are shown below in Table 13-17.

Ribozyme Cleavage of TERT RNA Target *in vitro*

Ribozymes targeted to the human TERT RNA are designed and synthesized as described above. These ribozymes can be tested for cleavage activity *in vitro*, for example using the following procedure. The target sequences and the nucleotide location within the TERT RNA are given in Tables 13-17.

Cleavage Reactions: Full-length or partially full-length, internally-labeled target RNA for ribozyme cleavage assay is prepared by *in vitro* transcription in the presence of [a-³²P] CTP, passed over a G 50 Sephadex column by spin chromatography and used as

substrate RNA without further purification. Alternately, substrates are 5'-³²P-end labeled using T4 polynucleotide kinase enzyme. Assays are performed by pre-warming 15 μ l of a 2X concentration of purified ribozyme in ribozyme cleavage buffer (50 mM Tris-HCl, pH 7.5 at 37°C, 10 mM MgCl₂) and the cleavage reaction was initiated by adding the 2X

5 ribozyme mix to an equal volume (15 μ l) of substrate RNA (maximum of 1-5 nM; 5×10^5 to 1×10^7 cpm) that was also pre-warmed in cleavage buffer. As an initial screen, assays are carried out for 1 hour at 37°C using a final concentration of either 40 nM or 1 mM ribozyme, *i.e.*, ribozyme excess. The reaction is quenched by the addition of an equal volume (30 μ l) of 95% formamide, 20 mM EDTA, 0.05% bromophenol blue and 0.05%

10 xylene cyanol after which the sample is heated to 95°C for 2 minutes, quick chilled and loaded onto a denaturing polyacrylamide gel. Substrate RNA and the specific RNA cleavage products generated by ribozyme cleavage are visualized on an autoradiograph of the gel. The percentage of cleavage is determined by Phosphor Imager[®] quantitation of bands representing the intact substrate and the cleavage products.

15

Example 2: PTP-1B

Protein tyrosine phosphorylation and dephosphorylation are important mechanisms in the regulation of signal transduction pathways that control the processes of cell growth, proliferation, and differentiation (Fantl, W. J., 1993, Annu. Rev. Biochem., 62, 453-481).

20 Cooperative enzyme classes regulate protein tyrosine phosphorylation and dephosphorylation events. These broad classes of enzymes consist of the protein tyrosine kinases (PTKs) and protein tyrosine phosphatases (PTPs). PTKs and PTPs can exist as both receptor-type transmembrane proteins and as cytoplasmic protein enzymes. Receptor tyrosine kinases propagate signal transduction events via extracellular receptor-ligand

25 interactions that result in the activation of the tyrosine kinase portion of the PTK in the cytoplasmic domain. Receptor-like transmembrane PTPs function through extracellular ligand binding that modulates dephosphorylation of intracellular phosphotyrosine proteins via cytoplasmic phosphatase domains. Cytoplasmic PTKs and PTPs exert enzymatic activity without receptor-mediated ligand interactions, however, phosphorylation can

30 regulate the activity of these enzymes.

Protein tyrosine phosphatase 1B, a cytoplasmic PTP, was the first PTP to be isolated in homogeneous form (Tonks, N. K., 1988, J. Biol. Chem., 263, 6722-6730), characterized (Tonks, N. K., 1988, J. Biol. Chem., 263, 6731-6737), and sequenced (Charbonneau, H., 1989, Biochemistry, 86, 5252-5256). Cytoplasmic and receptor-like PTPs both share a catalytic domain characterized by eleven conserved amino acids containing cysteine and arginine residues that are critical for phosphatase activity (Streuli, M., 1990, EMBO, 9, 2399-2407). A cysteine residue at position 215 is responsible for the covalent attachment of phosphate to the enzyme (Guan, K., 1991, J. Biol. Chem., 266, 17026-17030). The crystal structure of human PTP1B defined the phosphate binding site of the enzyme as a glycine rich cleft at the surface of the molecule with cysteine 215 positioned at the base of this cleft. The location of cysteine 215 and the shape of the cleft provide specificity of PTPase activity for tyrosine residues but not for serine or threonine residues (Barford, D., 1994, Science, 263, 1397-1404).

Receptor tyrosine kinase and protein tyrosine phosphatase localization plays a key role in the regulation of phosphotyrosine mediated signal transduction. PTP-1B activity and specificity against a panel of receptor tyrosine kinases demonstrated clear differences between substrates, suggesting that cellular compartmentalization is a determinant in defining the activity and function of the enzyme (Lammers, R., 1993, J. Biol. Chem., 268, 22456-22462). Experiments have indicated that PTP-1B is localized predominantly in the endoplasmic reticulum via its 35 amino acid carboxyterminal sequence. PTP-1B is also tightly associated with microsomal membranes with its catalytic phosphatase domain oriented towards the cytoplasm (Frangioni, J. V., 1992, Cell, 68, 545-560).

PTP-1B has been identified as a negative regulator of the insulin response. PTP-1B is widely expressed in insulin sensitive tissues (Goldstein, B. J., 1993, Receptor, 3, 1-15). Isolated PTP-1B dephosphorylates the insulin receptor *in vitro* (Tonks, N. K., 1988, J. Biol. Chem., 263, 6731-6737). PTP-1B dephosphorylation of multiple phosphotyrosine residues of the insulin receptor proceeds sequentially and with specificity for the three tyrosine residues that are critical for receptor autoactivation (Ramachandran, C., 1992, Biochemistry, 31, 4232-4238). In addition to insulin receptor dephosphorylation, PTP-1B also dephosphorylates the insulin related substrate 1 (IRS-1), a principal substrate of the insulin receptor (Lammers, R., 1993, J. Biol. Chem., 268, 22456-22462).

Microinjection of PTP1B into *Xenopus* oocytes results in the inhibition of insulin stimulated tyrosine phosphorylation of endogenous proteins, including the β -subunit of the insulin and insulin-like growth factor receptor proteins. The resulting 3 to 5 fold increase over endogenous PTPase activity also blocks the activation of an S6 peptide kinase (Cicirelli, M. F., 1990, Proc. Natl. Acad. Sci., 87, 5514-5518). Inactivation of recombinant rat PTP-1B with antibody immunoprecipitation results in the dramatic increase in insulin stimulated DNA synthesis and phosphatidylinositol 3'-kinase activity. Insulin stimulated receptor autophosphorylation and insulin receptor substrate 1 tyrosine phosphorylation are increased dramatically as well through PTP-1B inhibition (Ahmad, F., 1995, J. Biol. Chem., 270, 20503-20508).

Increased PTP-1B expression correlates with insulin resistance in hyperglycemic cultured fibroblasts. In this study, desensitized insulin receptor function was observed via impaired insulin-induced autophosphorylation of the receptor. Treatment with insulin sensitivity normalizing thiazolidine derivatives resulted in the amelioration of the hyperglycemic insulin resistance via a normalization in PTP-1B expression (Maegawa, H., 1995, J. Biol. Chem., 270, 7724-7730). A murine model of insulin resistance with a knockout of the heterotrimeric GTP-binding protein subunit $G_{i\alpha 2}$ provides a type 2 diabetes phenotype that correlates with the increased expression of PTP-1B (Moxam, C. M., 1996, Nature, 379, 840-844).

PTP-1B interacts directly with the activated insulin receptor β -subunit. An inactive homolog of PTP-1B was used to precipitate the activated insulin receptor in both purified receptor preparations and whole-cell lysates. Phosphorylation of the insulin receptor's triple tyrosine residues in the kinase domain is necessary for PTP-1B interaction. Furthermore, insulin stimulates tyrosine phosphorylation of PTP-1B (Seely, B. L., 1996, Diabetes, 45, 1379-1385). A similar study confirmed the direct interaction of PTP-1B with the insulin receptor β -subunit as well as the required multiple phosphorylation sites within the receptor and PTP-1B (Bandyopadhyay, D., J. Biol. Chem., 272, 1639-1645).

Knockout mice lacking the PTP-1B gene (both homozygous PTP-1B^{-/-} and heterozygous PTP-1B^{+/-}) have been used to study the specific role of PTP-1B relating to insulin action *in vivo*. The resulting PTP-1B deficient mice were healthy and, in the fed state, had lower blood glucose and circulating insulin levels that were half that of their

PTP-1B^{+/+} expressing littermates. These PTP-1B deficient mice demonstrated enhanced insulin sensitivity in glucose and insulin tolerance tests. At the physiological level, the PTP-1B deficient mice showed increased phosphorylation of the insulin receptor after insulin administration. When fed a high fat diet, the PTP-1B deficient mice were resistant to weight gain and remained insulin sensitive as opposed to normal PTP-1B expressing mice, who rapidly gained weight and become insulin resistant (Elchebly, M., 1999, Science, 283, 1544-1548). As such, modulation of PTP-1B expression could be used to regulate autophosphorylation of the insulin receptor and increase insulin sensitivity *in vivo*. This modulation could prove beneficial in the treatment of insulin related disease states.

10 In light of the above findings, particular disease states that involve PTP-1B expression include but are not limited to:

Diabetes: Both type 1 and type 2 diabetes may be treated by modulation of PTP-1B expression. Type 2 diabetes correlates to desensitized insulin receptor function (White *et al.*, 1994). Disruption of the PTP-1B dephosphorylation of the insulin receptor *in vivo* manifests in insulin sensitivity and increased insulin receptor autophosphorylation (Elchebly *et al.*, 1999). Insulin dependant diabetes, type 1, may respond to PTP-1B modulation through increased insulin sensitivity.

Obesity: Elchebly *et al.*, 1999, demonstrated that PTP-1B deficient mice were resistant to weight gain when fed a high fat diet compared to normal PTP-1B expressing mice. This finding suggests that PTP-1B modulation may be beneficial in the treatment of obesity. Ahmad *et al.*, 1997, Metab. Clin. Exp., 46, 1140-1145, describe reduced PTPs in adipose tissue and improved insulin sensitivity in obese subjects following weight loss.

25 Troglitazone is a non-limiting example of a pharmaceutical agent that can be combined with or used in conjunction with the nucleic acid molecules (e.g. ribozymes and antisense molecules) of the instant invention. Those skilled in the art will recognize that other drugs such as anti-diabetes and anti-obesity compounds and therapies can be similarly be readily combined with the nucleic acid molecules of the instant invention (e.g. ribozymes and antisense molecules) are hence within the scope of the instant invention.

30 Methods have been developed to assay PTP-1B activity.

Maegawa *et al.*, 1995, J. Biol. Chem., 270, 7724-7730, describe a tissue culture model in which Rat 1 fibroblasts expressing human insulin receptors can be used to model

hyperglycemia induced insulin resistance. Maegawa *et al.* also describe assays to measure PTPase activity using labeled phosphorylated insulin receptors and by immunoenzymatic techniques.

- 5 Moxham *et al.*, 1996, Nature, 379, 840-844, describe a murine animal and tissue culture model employing *Gia2* deficiency to study hyperinsulinaemia, impaired glucose tolerance and resistance to insulin *in vivo*. Assays for PTPase activity and tyrosine phosphorylation of insulin-receptor substrate 1 are described.

- 10 Khandelwal *et al.*, 1995, Molecular and Cellular Biochemistry, 153, 87-94, describe four different animal models for studying insulin dependent and insulin resistant diabetes mellitus. These models were used to study the effect of vanadate, an insulin mimetic and PTPase inhibitor, on the insulin-stimulated phosphorylation of the insulin receptor and its tyrosine kinase activity.

Wang *et al.*, 1999, Biochim. Biophys. Acta, 1431, 14-23, describe fluorescein monophosphates as fluorogenic substrates for PTPs.

- 15 Various methods and compounds have been developed to inhibit protein tyrosine phosphatase activity.

Wrobel *et al.*, 1999, J. Med. Chem., 42, 3199-3202, describe PTP-1B inhibition and antihyperglycemic activity in the ob/ob mouse model by 11-arylbenzo[b]naphtho[2,3-d]furans and arylbenzo[b]naphtho[2,3-d]thiophenes.

- 20 Andersen *et al.*, International PCT publication No. WO 98/DK407 describe the preparation of thienopyridzinones and thienochromenones as modulators of PTPases.

Taing *et al.*, 1999, Biochemistry, 38, 3793-3803, describe potent and highly selective inhibitors of PTP-1B comprising an array of bis(aryldifluorophosphonates).

- 25 Ham *et al.*, 1999, Bioorg. Med. Chem. Lett., 9, 185-186, describe selective inactivation of PTP-1B by a sulfone analog of naphthoquinone.

Desmarais *et al.*, 1999, Biochem. J., 337, 219-223, describe [Difluoro(phosphono)methyl]phenylalanine-containing peptide inhibitors of PTPs.

Taylor *et al.*, 1998, Bioorg. Med. Chem., 6, 2235, describe potent non-peptidyl inhibitors of PTP-1B.

- 30 Kotoris *et al.*, 1998, Bioorg. Med. Chem. Lett., 8, 3275-3280, describe novel phosphate mimetics for the design of non-peptidyl inhibitors of PTPs.

Groves *et al.*, 1998, *Biochemistry*, 37, 17773-17783, describe the structural basis for PTP-1B inhibition by the phosphotyrosine peptide mimetics (difluoronaphthylmethyl)phosphonic acid and the fluoromalonyl tyrosines with complexed crystal structures.

- 5 Yao *et al.*, 1998, *Bioorgl Med. Chem.*, 6, 1799-1810, describe the structure-based design and synthesis of small molecule PTP-1B inhibitors comprising novel naphthyldifluoromethyl phosphonic acids 1 and 2.

Taylor *et al.*, 1998, *Bioorg. Med. Chem.*, 6, 1457-1468, describe potent non-peptidyl inhibitors of PTP-1B.

- 10 Desmarais *et al.*, 1998, *Arch. Biochem. Biophys.*, 354, 225-231, describe inhibition of PTP-1B and CD45 by sulfotyrosyl peptides.

Mjalli *et al.*, application US 96-766114, cont. in part of US patent No. 543,630, describe the preparation of heterocyclic compounds as modulators of proteins with phosphotyrosine recognition units.

- 15 Wang *et al.*, 1998, *Bioorg. Med. Chem. Lett.*, 8, 345-350, describe naphthalenebis[α,α -difluoromethylenephosphonates] as potent inhibitors of PTPs.

Rice *et al.*, 1997, *Biochemistry*, 36, 15965-15974, describe a targeted library of small molecule tyrosine and dual-specificity phosphatase inhibitors with random side chain variation from a rational core design.

- 20 Olefsky, International PCT publication No. WO 97/US2752 describes a method and phosphopeptides used for the treatment of insulin resistance based on the association of PTP-1B with the activated insulin receptor. Also included is a method for determining whether a compound inhibits PTP-1B binding to the insulin receptor.

- 25 Huyer *et al.*, 1997, *J. Biol. Chem.*, 272, 843-851, describe the mechanism of inhibition of PTPases by vanadate and pervanadate.

Burke *et al.*, 1996, *Biochemistry*, 35, 15989-15996, describe the structure-based design of PTP-1B inhibitors.

- 30 Tonks *et al.*, International PCT publication No. WO 97/US13016, describe substrate-trapping protein PTPase mutants for identification of tyrosine-phosphorylated protein substrates and their clinical uses.

The human genome is thought to contain up to 100 PTPases, each varying slightly in chemistry but vastly in function. Compounds designed to inhibit PTP-1B activity specifically by covalent binding to or modification of PTP-1B have the potential for multiple side effects. Conventional drug substances that will potently suppress PTP-1B activity with few or no side effects from interaction with other PTPs are difficult to envision. A more attractive approach to PTP-1B modulation would involve the specific regulation of PTP-1B expression with oligonucleotides.

Identification of Potential Target Sites in Human PTP-1B RNA

The sequence of human PTP-1B was screened for accessible sites using a computer folding algorithm. Regions of the RNA that did not form secondary folding structures and contained potential ribozyme and/or antisense binding/cleavage sites were identified. The sequences of these cleavage sites are shown in Tables 3-8.

Selection of Enzymatic Nucleic Acid Cleavage Sites in Human PTP-1B RNA

To test whether the sites predicted by the computer-based RNA folding algorithm corresponded to accessible sites in PTP-1B RNA, 10 hammerhead ribozyme, five NCH and three G-Cleaver ribozyme sites were selected for further analysis (Table 8). Ribozyme target sites were chosen by analyzing sequences of Human PTP-1B (Genbank accession number M33689) and prioritizing the sites on the basis of folding. Ribozymes were designed that could bind each target and were individually analyzed by computer folding (Christoffersen *et al.*, 1994 *J. Mol. Struc. Theochem*, 311, 273; Jaeger *et al.*, 1989, *Proc. Natl. Acad. Sci. USA*, 86, 7706) to assess whether the ribozyme sequences fold into the appropriate secondary structure. Those ribozymes with unfavorable intramolecular interactions between the binding arms and the catalytic core were eliminated from consideration. As noted below, varying binding arm lengths can be chosen to optimize activity. Generally, at least 5 bases on each arm are able to bind to, or otherwise interact with, the target RNA.

Chemical Synthesis and Purification of Ribozymes for Efficient Cleavage of PTP-1B RNA

Ribozymes were designed to anneal to various sites in the RNA message. The binding arms are complementary to the target site sequences described above. The ribozymes were chemically synthesized. The method of synthesis used followed the procedure for normal RNA synthesis as described above and in Usman et al., (1987 J. Am. Chem. Soc., 109, 7845), Scaringe et al., (1990 Nucleic Acids Res., 18, 5433) and Wincott et al., supra, and made use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. The average stepwise coupling yields were >98%.

Ribozymes were also synthesized from DNA templates using bacteriophage T7 RNA polymerase (Milligan and Uhlenbeck, 1989, Methods Enzymol. 180, 51). Ribozymes were purified by gel electrophoresis using general methods or were purified by high pressure liquid chromatography (HPLC; see Wincott et al., supra; the totality of which is hereby incorporated herein by reference) and were resuspended in water. The sequences of the chemically synthesized ribozymes used in this study are shown below in Tables 3-8.

Ribozyme Cleavage of PTP-1B RNA Target *in vitro*

Ribozymes targeted to the human PTP-1B RNA are designed and synthesized as described above. These ribozymes can be tested for cleavage activity *in vitro*, for example, using the following procedure. The target sequences and the nucleotide location within the PTP-1B RNA are given in Tables 3-8.

Cleavage Reactions: Full-length or partially full-length, internally-labeled target RNA for ribozyme cleavage assay is prepared by *in vitro* transcription in the presence of [α -³²P] CTP, passed over a G 50 Sephadex column by spin chromatography and used as substrate RNA without further purification. Alternately, substrates are 5'-³²P-end labeled using T4 polynucleotide kinase enzyme. Assays are performed by pre-warming a 2X concentration of purified ribozyme in ribozyme cleavage buffer (50 mM Tris-HCl, pH 7.5 at 37°C, 10 mM MgCl₂) and the cleavage reaction was initiated by adding the 2X ribozyme mix to an equal volume of substrate RNA (maximum of 1-5 nM) that was also pre-warmed in cleavage buffer. As an initial screen, assays are carried out for 1 hour at 37°C using a final concentration of either 40 nM or 1 mM ribozyme, *i.e.*, ribozyme excess.

The reaction is quenched by the addition of an equal volume of 95% formamide, 20 mM EDTA, 0.05% bromophenol blue and 0.05% xylene cyanol after which the sample is heated to 95°C for 2 minutes, quick chilled and loaded onto a denaturing polyacrylamide gel. Substrate RNA and the specific RNA cleavage products generated by ribozyme
5 cleavage are visualized on an autoradiograph of the gel. The percentage of cleavage is determined by Phosphor Imager® quantitation of bands representing the intact substrate and the cleavage products.

Example 3: MetAP-2

10 Methionyl aminopeptidases are metalloproteases that are known to possess post-translational enzymatic activity by hydrolytically cleaving amino-terminal methionine residues from nascent peptide substrates in a non-processive manner (Kendall, R. L., 1992, J. Biol. Chem., 267, 20667-20673). This family of enzymes is divided into two classes (type 1 and type 2) based on differences in sequence, although the overall structure of the
15 two classes are similar (Liu, S., 1998, Science, 282, 1324-1327). Methionine aminopeptidase expression appears to be involved in the control of cellular proliferation. Deletion of the MetAP gene from *E. Coli* is lethal (Chang, S. Y., 1989, J. Bacteriol., 171, 4071-4072). In *Saccharomyces cerevisiae*, deletion of the gene that codes for either MetAP-1 or 2 results in a slow growth phenotype while deletion of both genes is lethal (Li,
20 X., 1995, Proc. Natl. Acad. Sci., 92, 12357-12361). (Human methionine aminopeptidase-1, MetAP-1, accession No. P53582).

The aminopeptidase function of this class of enzymes may serve a regulatory role in activating signal peptides in conjunction with N-myristoyl transferase (NMT) activity. NMT is expressed from a lethal gene in yeast (Duronio, R. J., 1989, Science, 243, 796-
25 800). NMT is responsible for amino-terminal ligation of myristic acid onto nascent peptides and cannot act on peptides with an amino-terminal methionine residue (Resh, M. D., 1996, Cell. Signal., 8, 403-412). Myristoylation of proteins correlates to intracellular localization events that may determine why certain signaling proteins are dependent on NMT for activity (Taunton, J., 1997, Chemistry & Biology, 4, 493-496). Protein tyrosine
30 kinase Src is dependant on myristoylation for activity and has been identified as an upstream regulator of human vascular endothelial growth factor (VEGF) expression

through hypoxic induction in solid tumors (Mukhopadhyay, D., 1995, Nature, 375, 577-581). MetAPs may therefore regulate the activation of signal peptides (such as VEGF) through cotranslational modification of nascent peptides with NMT. Disruption of protein myristoylation by MetAP inhibition could result in the improper localization of signaling proteins resulting in inhibition of cell growth. (Human N-myristoyltransferase, hNMT, 5 accession No. AF043324.)

Fumagillin, a sesquiterpene diepoxide metabolite of the fungus *Aspergillus fumigatus*, and a related compound TNP-470, are strong inhibitors of growth in cultured endothelial cells. The antiproliferative and angiostatic activity of fumagillin was originally 10 discovered by the serendipitous contamination of *Aspergillus fumigatus* in an endothelial cell culture dish in which cells closest to the fungal colony displayed growth inhibition. Synthetic analogs of fumagillin were later synthesized resulting in the discovery of TNP-470, which is 50 times more potent of an inhibitor than fumagillin and is less toxic in mice (Ingber, D., 1990, Nature, 348, 555-557). Treatment of endothelial cells with these 15 compounds results in late G1 phase arrest. TNP-470 inhibits the signaling pathway of retinoblastoma gene product phosphorylation, cyclin dependent kinases cdk2 and cdk4 activation, and cyclins E and A expression (Abe, J., 1994, Cancer Res., 54, 3407-3412). TNP-470 has also been shown to potently inhibit endothelial cell proliferation induced by the growth factors VEGF and bFGF (Toi, M., 1994, Oncology Reports, 1, 423-426).

20 The bifunctional protein MetAP-2 has been identified as the molecular target for fumagillin and related compounds that demonstrate antiproliferative activity in endothelial cells. The use of affinity chromatography with a fumagillin-biotin conjugate resulted in the isolation of a 67-kDa mammalian protein through covalent interaction with the bound substrate. Analysis of digested peptide fragments from the isolated protein revealed 25 MetAP-2 as the covalently bound substrate. Subsequent growth inhibition studies in yeast utilizing MetAP-1 and MetAP-2 deletion strains determined that MetAP-2 is selectively inhibited by fumagillin *in vivo* (Sin, N., 1997, Proc. Natl. Acad. Sci., 94, 6099-6103). A similar study with TNP-470 and ovalicin, another potent inhibitor of neovascularization, determined that MetAP-2 is the molecular target for these fumagillin-related compounds 30 (Griffith, E. C., 1997, Chemistry & Biology, 4, 461-471).

MetAP-2 expression correlates with cellular growth. Non-dividing cells in culture have no detectable levels of the 67-kDa MetAP-2 protein by immunoassay. MetAP-2 has been shown to affect translational initiation by association with eukaryotic initiation factor 2 α (eIF-2 α) (Ray, M. K., 1992, Proc. Natl. Acad. Sci., 89, 539-543). The binding of
5 MetAP-2 with eIF-2 α inhibits the heme-regulated inhibitor kinase (HRI) phosphorylation of eIF-2 α *in vitro* in reticulocyte lysates (Datta, B., 1988, Proc. Natl. Acad. Sci., 85, 3324-3328). MetAP-2/eIF-2 α binding results in the partial reversal of protein synthesis inhibition by double stranded RNA dependent kinase mediated phosphorylation *in vivo* (Wu, S., 1996, Biochemistry, 35, 8275-8280). Griffith *et al.* also determined that covalent
10 binding of TNP-470 and ovalicin, while potently inhibiting methionine aminopeptidase type 2 activity specifically, did not affect the regulatory activity of MetAP-2 on eIF-2 α . This finding by Griffith *et al.* rules out the possibility that control of eIF-2 α phosphorylation by MetAP-2 is responsible for the inhibition of endothelial cell proliferation by fumagillin related compounds.

15 Particular angiogenesis related degenerative and disease states that can be associated with MetAP expression modulation include but are not limited to:

Cancer: Solid tumors are unable to grow or metastasize without the formation of new blood vessels (Hanahan, D., 1996, Cell, 86, 353-364). Inhibition of angiogenesis via MetAP modulation can potentially be used to treat a wide variety of cancers.

20

Diabetic retinopathy and age related macular degeneration: Ocular neovascularization is observed in diabetic retinopathy, which is mediated by up-regulation of VEGF (Adamis, A. P., 1994, Amer. J. Ophthal., 118, 445-450). The requirement of protein kinase Src in hypoxia induced VEGF expression (Mukhopadhyay, D., 1995, Nature, 375, 577-581) indicates that MetAP modulation of aminopeptidase activity can
25 potentially be used to treat conditions involving ocular neovascularization.

Arthritis: The ingrowth of a vascular pannus in arthritis may be mediated by the overexpression of angiogenic factors from infiltrating inflammatory cells, macrophages, and immune cells (Peacock, D. J., 1992, J. exp. Med., 175, 1135-1138). Angiogenesis
30 inhibition through MetAP modulation can potentially be used to treat arthritis.

Psoriasis: Angiogenesis has been implicated in psoriasis due to overexpression of the angiogenic polypeptide interleukin-8 and decreased expression of the angiogenesis inhibitor thrombospondin (Nickoloff, B. J., 1994, Amer. J. Pathol. 44, 820-828).

Angiogenesis inhibition through MetAP modulation can potentially be used to treat
5 psoriasis.

Female reproduction: Angiogenesis in the female reproductive system has been implicated in several disorders of the reproductive tract (Reynolds, L. P., 1992, FASEB, 6, 886-892). Modulation of angiogenesis through control of MetAP may have various applications in the area of female reproduction and fertility.

10 Various methods have been developed to assay MetAP activity.

Griffith *et al.*, 1998, Proc. Natl. Acad. Sci., 95, 15183-15188, describe an enzymatic assay for MetAP-2 activity *in vitro* and an endothelial cell culture proliferation assay for MetAP-2 activity *in vivo*.

Weber *et al.*, 1999, International PCT publication No. WO 98/US-21231 describe
15 novel fluorescent reporter molecules and an enzymatic assay that can be used for determining the activity of MetAP-2 for drug screening and determining the chemosensitivity of human cancer cells to treatment with chemotherapeutic drugs.

Larrabee, J. A. *et al.*, 1999, Anal. Biochem, 269, 194-198, describe the use of a high-pressure liquid chromatographic (HPLC) method for assaying MetAP-2 activity with
20 application to the study of enzymic inactivation.

Quantitative methods have been developed to assay the efficacy of antiangiogenic therapies.

Wantanabe *et al.*, 1992, Molec. Biol. Cell, 3, 324a, describe the quantitation of angiogenic peptides (bFGF) in human serum as a prognostic test for breast cancer.

25 Nguyen *et al.*, 1994, J. Natn. Cancer Inst., 86, 356-361, describe the quantitation of angiogenic peptides (bFGF) in the urine of patients with a wide spectrum of cancers.

Li *et al.*, 1994, The Lancet, 344, 82-86, describe the quantitation of angiogenic peptides (bFGF) in the cerebrospinal fluid of children with brain tumors. This work also describes determining the extent of neovascularization in histological sections by utilizing
30 microvessel count.

The present body of knowledge in angiogenesis research indicates the need for compounds that can modulate MetAP activity for research, diagnostic, trait alteration, animal health and therapeutic use.

Griffith *et al.*, International PCT publication No. WO 9856372 describe small molecule inhibitors of MetAP2 and uses thereof.

D'Amato *et al.*, International PCT publication No. WO 9805293 describe the use of AGM-1470 (TNP-470) as an angiogenesis inhibitor for use in regulating the female reproductive system and for treating diseases of the reproductive tissue.

Davidson *et al.*, US patent No. 5,801,146 describe a compound and method for inhibiting angiogenesis using mammalian krigle 5 protein.

Cao *et al.*, US patent No. 5,854,221 describe a protein-based endothelial cell proliferation inhibitor and its method of use.

Chang *et al.*, US patent No. 5,888,796 describe a clone of a nucleotide sequence encoding a protein having two functions comprising methionine aminopeptidase activity and anti eIF-2 phosphorylation activity.

Wang *et al.*, 1998, Proc. Am. Assoc. Cancer Res., 39, 98 (abstr.) describe blocked proliferation of human endothelial cells by human MetAP-2 antisense oligonucleotides.

A rat corneal model has been developed to study ribozyme inhibition of VEGF receptor-mediated angiogenesis (Pavco, P. A., 1999, Nucleic Acids Research, 27, 2569-2577). A similar study employing MetAP-2 inhibition could be used to study ribozyme based inhibition of MetAP-2 induced angiogenesis *in vivo*.

Identification of Potential Target Sites in Human MetAP-2 RNA

The sequence of human MetAP-2 was screened for accessible sites using a computer-folding algorithm. Regions of the RNA that did not form secondary folding structures and contained potential ribozyme and/or antisense binding/cleavage sites were identified. The sequences of these cleavage sites are shown in Tables 9-12.

Selection of Enzymatic Nucleic Acid Cleavage Sites in Human MetAP-2 RNA

To test whether the sites predicted by the computer-based RNA folding algorithm corresponded to accessible sites in MetAP-2 RNA, 11 hammerhead ribozyme, 4 NCH and three G-Cleaver ribozyme sites were selected for further analysis (Table 12). Ribozyme

target sites were chosen by analyzing sequences of Human MetAP-2 (Genbank accession number HSU29607) and prioritizing the sites on the basis of folding. Ribozymes were designed that could bind each target and were individually analyzed by computer folding (Christoffersen *et al.*, 1994 *J. Mol. Struc. Theochem*, 311, 273; Jaeger *et al.*, 1989, *Proc. Natl. Acad. Sci. USA*, 86, 7706) to assess whether the ribozyme sequences fold into the appropriate secondary structure. Those ribozymes with unfavorable intramolecular interactions between the binding arms and the catalytic core were eliminated from consideration. As noted below, varying binding arm lengths can be chosen to optimize activity. Generally, at least 5 bases on each arm are able to bind to, or otherwise interact with, the target RNA.

Chemical Synthesis and Purification of Ribozymes for Efficient Cleavage of MetAP-2 RNA

Ribozymes were designed to anneal to various sites in the RNA message. The binding arms are complementary to the target site sequences described above. The ribozymes were chemically synthesized. The method of synthesis used followed the procedure for normal RNA synthesis as described above and in Usman *et al.*, (1987 *J. Am. Chem. Soc.*, 109, 7845), Scaringe *et al.*, (1990 *Nucleic Acids Res.*, 18, 5433) and Wincott *et al.*, *supra*, and made use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. The average stepwise coupling yields were >98%.

Ribozymes were also synthesized from DNA templates using bacteriophage T7 RNA polymerase (Milligan and Uhlenbeck, 1989, *Methods Enzymol.* 180, 51). Ribozymes were purified by gel electrophoresis using general methods or were purified by high pressure liquid chromatography (HPLC; see Wincott *et al.*, *supra*; the totality of which is hereby incorporated herein by reference) and were resuspended in water. The sequences of the chemically synthesized ribozymes used in this study are shown below in Table 9-12.

Ribozyme Cleavage of MetAP-2 RNA Target *in vitro*

Ribozymes targeted to the human MetAP-2 RNA are designed and synthesized as described above. These ribozymes can be tested for cleavage activity *in vitro*, for

example, using the following procedure. The target sequences and the nucleotide location within the MetAP-2 RNA are given in Tables 9-12.

Cleavage Reactions: Full-length or partially full-length, internally-labeled target RNA for ribozyme cleavage assay is prepared by *in vitro* transcription in the presence of [a-³²P] CTP, passed over a G 50 Sephadex column by spin chromatography and used as substrate RNA without further purification. Alternately, substrates are 5'-³²P-end labeled using T4 polynucleotide kinase enzyme. Assays are performed by pre-warming a 2X concentration of purified ribozyme in ribozyme cleavage buffer (50 mM Tris-HCl, pH 7.5 at 37°C, 10 mM MgCl₂) and the cleavage reaction was initiated by adding the 2X ribozyme mix to an equal volume of substrate RNA (maximum of 1-5 nM) that was also pre-warmed in cleavage buffer. As an initial screen, assays are carried out for 1 hour at 37°C using a final concentration of either 40 nM or 1 mM ribozyme, *i.e.*, ribozyme excess. The reaction is quenched by the addition of an equal volume of 95% formamide, 20 mM EDTA, 0.05% bromophenol blue and 0.05% xylene cyanol after which the sample is heated to 95°C for 2 minutes, quick chilled and loaded onto a denaturing polyacrylamide gel. Substrate RNA and the specific RNA cleavage products generated by ribozyme cleavage are visualized on an autoradiograph of the gel. The percentage of cleavage is determined by Phosphor Imager[®] quantitation of bands representing the intact substrate and the cleavage products.

Example 4: BACE, ps-1, ps-2

Alzheimer's disease (AD) is a progressive, degenerative disease of the brain which affects approximately 4 million people in the United States alone. An estimated 14 million Americans will have Alzheimer's disease by the middle of the next century if no cure or definitive prevention of the disease is found. Nearly one out of ten people over age 65 and nearly half of those over 85 have Alzheimer's disease. Alzheimer's disease is not confined to the elderly, a small percentage of people in their 30's and 40's are afflicted with early onset AD. Alzheimer's disease is the most common form of dementia, and amounts to the third most expensive disease in the US following heart disease and cancer. An estimated 100 billion dollars are spent annually on Alzheimer's disease (National Alzheimer's Association, 1999).

Alzheimer's disease is characterized by the progressive formation of insoluble plaques and vascular deposits in the brain consisting of the 4 kD amyloid β peptide ($A\beta$). These plaques are characterized by dystrophic neurites that show profound synaptic loss, neurofibrillary tangle formation, and gliosis. $A\beta$ arises from the proteolytic cleavage of the large type I transmembrane protein, β -amyloid precursor protein (APP) (Kang *et al.*, 1987, *Nature*, 325, 733). Processing of APP to generate $A\beta$ requires two sites of cleavage by a β -secretase and a γ -secretase. β -secretase cleavage of APP results in the cytoplasmic release of a 100 kD soluble amino-terminal fragment, APPs β , leaving behind a 12 kD transmembrane carboxy-terminal fragment, C99. Alternately, APP can be cleaved by a α -secretase to generate cytoplasmic APPs α and transmembrane C83 fragments. Both remaining transmembrane fragments, C99 and C83, can be further cleaved by a γ -secretase, leading to the release and secretion of Alzheimer's related $A\beta$ and a non-pathogenic peptide, p3, respectively (Vassar *et al.*, 1999, *Science*, 286, 735-741). Early onset familial Alzheimer's disease is characterized by mutant APP protein with a Met to Leu substitution at position P1, characterized as the "Swedish" familial mutation (Mullan *et al.*, 1992, *Nature Genet.*, 1, 345). This APP mutation is characterized by a dramatic enhancement in β -secretase cleavage (Citron *et al.*, 1992, *Nature*, 360, 672).

The identification of β -secretase, and γ -secretase constituents involved in the release of β -amyloid protein is of primary importance in the development of treatment strategies for Alzheimer's disease. Characterization of α -secretase is also important in this regard since α -secretase cleavage may compete with β -secretase cleavage resulting in non-pathogenic vs. pathogenic protein production. Involvement of the two metalloproteases, ADAM 10, and TACE has been demonstrated in α -cleavage of APP (Buxbaum *et al.*, 1999, *J. Biol. Chem.*, 273, 27765, and Lammich *et al.*, 1999, *Proc. Natl. Acad. Sci. U.S.A.*, 96, 3922). Studies of γ -secretase activity have demonstrated presenilin dependence (De Strooper *et al.*, 1998, *Nature*, 391, 387, and De Strooper *et al.*, 1999, *Nature*, 398, 518), and as such, presenilins have been proposed as γ -secretase even though presenilin does not present proteolytic activity (Wolfe *et al.*, 1999, *Nature*, 398, 513).

Recently, Vassar *et al.*, 1999, *supra* reported β -secretase cleavage of AAP by the transmembrane aspartic protease beta site APP cleaving enzyme, BACE. While other potential candidates for β -secretase have been proposed (for review see Evin *et al.*, 1999, *Proc. Natl. Acad. Sci. U.S.A.*, 96, 3922), none have demonstrated the full range of characteristics expected from this enzyme. Vassar *et al, supra*, demonstrate that BACE expression and localization are as expected for β -secretase, that BACE overexpression in cells results in increased β -secretase cleavage of APP and Swedish APP, that isolated BACE demonstrates site specific proteolytic activity on APP derived peptide substrates, and that antisense mediated endogenous BACE inhibition results in dramatically reduced β -secretase activity.

Current treatment strategies for Alzheimer's disease rely on either the prevention or the alleviation of symptoms and/or the slowing down of disease progression. Two drugs approved in the treatment of Alzheimer's, donepezil (Aricept®) and tacrine (Cognex®), both cholinomimetics, attempt to slow the loss of cognitive ability by increasing the amount of acetylcholine available to the brain. Antioxidant therapy through the use of antioxidant compounds such as alpha-tocopherol (vitamin E), melatonin, and selegiline (Eldepryl®) attempt to slow disease progression by minimizing free radical damage. Estrogen replacement therapy is thought to incur a possible preventative benefit in the development of Alzheimer's disease based on limited data. The use of anti-inflammatory drugs may be associated with a reduced risk of Alzheimer's as well. Calcium channel blockers such as Nimodipine® are considered to have a potential benefit in treating Alzheimer's disease due to protection of nerve cells from calcium overload, thereby prolonging nerve cell survival. Nootropic compounds, such as acetyl-L-carnitine (Alcar®) and insulin, have been proposed to have some benefit in treating Alzheimer's due to enhancement of cognitive and memory function based on cellular metabolism.

Whereby the above treatment strategies may all improve quality of life in Alzheimer's patients, there exists an unmet need in the comprehensive treatment and prevention of this disease. As such, there exists the need for therapeutics effective in reversing the physiological changes associated with Alzheimer's disease, specifically, therapeutics that can eliminate and/or reverse the deposition of amyloid β peptide. The use of compounds to modulate the expression of proteases that are instrumental in the

release of amyloid β peptide, namely β -secretase (BACE), and γ -secretase (presenilin), is of therapeutic significance.

5 Tsai *et al.*, 1999, Book of Abstracts, 218th ACS National Meeting, New Orleans, Aug 22-26, describe substrate-based alpha-aminoisobutyric acid derivatives of difluoro ketone peptidomimetic inhibitors of amyloid β peptide through γ -secretase inhibition.

Czech *et al.*, International PCT publication No. WO/9921886, describe peptides capable of inhibiting the interaction between presenilins and the β -amyloid peptide or its precursor for therapeutic use.

10 Fournier *et al.*, International PCT publication No. WO/9916874, describe human brain proteins capable of interacting with presenilins and cDNAs encoding them toward therapeutic use.

St. George-Hyslop *et al.*, International PCT publication No. WO/9727296, describe genes for proteins that interact with presenilins and their role in Alzheimer's disease toward therapeutic use.

15 Vassar *et al.*, 1999, *Science*, 286, 735-741, describe specific antisense oligonucleotides targeting BACE, used for inhibition studies of endogenous BACE expression in 101 cells and APPsw cells via lipid mediated transfection.

20 Vassar *et al.*, 1999, *Science*, 286, 735-741, describe a cell culture model for studying BACE inhibition. Specific antisense nucleic acid molecules targeting BACE mRNA were used for inhibition studies of endogenous BACE expression in 101 cells and APPsw (Swedish type amyloid precursor protein expressing) cells via lipid mediated transfection. Antisense treatment resulted in dramatic reduction of both BACE mRNA by Northern blot analysis, and APPs β sw ("Swedish" type β -secretase cleavage product) by ELISA, with maximum inhibition of both parameters at 75-80%. This model was also used to study the effect of BACE inhibition on amyloid β -peptide production in APPsw cells.

25 Games *et al.*, 1995, *Nature*, 373, 523-527, describe a transgenic mouse model in which mutant human familial type APP (Phe 717 instead of Val) is overexpressed. This model results in mice that progressively develop many of the pathological hallmarks of Alzheimer's disease, and as such, provides a model for testing therapeutic drugs.

30 Particular degenerative and disease states that can be associated with BACE expression modulation include but are not limited to Alzheimer's disease and dementia.

Donepezil, tacrine, selegeline, and acetyl-L-carnitine are non-limiting examples of pharmaceutical agents that can be combined with or used in conjunction with the nucleic acid molecules (e.g. ribozymes and antisense molecules) of the instant invention. Those skilled in the art will recognize that other drugs such as diuretic and antihypertensive compounds and therapies can be similarly be readily combined with the nucleic acid molecules of the instant invention (e.g. ribozymes and antisense molecules) are hence within the scope of the instant invention.

Identification of Potential Target Sites in Human BACE RNA

The sequence of human BACE was screened for accessible sites using a computer-folding algorithm. Regions of the RNA that did not form secondary folding structures and contained potential ribozyme and/or antisense binding/cleavage sites were identified. The sequences of these cleavage sites are shown in Tables 18-23.

Selection of Enzymatic Nucleic Acid Cleavage Sites in Human BACE RNA

Ribozyme target sites were chosen by analyzing sequences of Human BACE (Genbank sequence accession number: AF190725) and prioritizing the sites on the basis of folding. Ribozymes were designed that could bind each target and were individually analyzed by computer folding (Christoffersen *et al.*, 1994 *J. Mol. Struct. Theochem*, 311, 273; Jaeger *et al.*, 1989, *Proc. Natl. Acad. Sci. USA*, 86, 7706) to assess whether the ribozyme sequences fold into the appropriate secondary structure. Those ribozymes with unfavorable intramolecular interactions between the binding arms and the catalytic core were eliminated from consideration. As noted below, varying binding arm lengths can be chosen to optimize activity. Generally, at least 5 bases on each arm are able to bind to, or otherwise interact with, the target RNA.

Chemical Synthesis and Purification of Ribozymes and Antisense for Efficient Cleavage and/or blocking of BACE RNA

Ribozymes and antisense constructs were designed to anneal to various sites in the RNA message. The binding arms of the ribozymes are complementary to the target site sequences described above, while the antisense constructs are fully complimentary to the target site sequences described above. The ribozymes and antisense constructs were

chemically synthesized. The method of synthesis used followed the procedure for normal RNA synthesis as described above and in Usman et al., (1987 *J. Am. Chem. Soc.*, 109, 7845), Scaringe et al., (1990 *Nucleic Acids Res.*, 18, 5433) and Wincott et al., *supra*, and made use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. The average stepwise coupling yields were >98%.

Ribozymes and antisense constructs were also synthesized from DNA templates using bacteriophage T7 RNA polymerase (Milligan and Uhlenbeck, 1989, *Methods Enzymol.* 180, 51). Ribozymes and antisense constructs were purified by gel electrophoresis using general methods or were purified by high pressure liquid chromatography (HPLC; See Wincott et al., *supra*; the totality of which is hereby incorporated herein by reference) and were resuspended in water. The sequences of the chemically synthesized ribozymes and antisense constructs used in this study are shown below in **Table 18-23**.

15

Ribozyme Cleavage of BACE RNA Target *in vitro*

Ribozymes targeted to the human BACE RNA are designed and synthesized as described above. These ribozymes can be tested for cleavage activity *in vitro*, for example, using the following procedure. The target sequences and the nucleotide location within the BACE RNA are given in **Tables 18-23**.

20

Cleavage Reactions: Full-length or partially full-length, internally-labeled target RNA for ribozyme cleavage assay is prepared by *in vitro* transcription in the presence of [α - 32 P] CTP, passed over a G 50 Sephadex column by spin chromatography and used as substrate RNA without further purification. Alternately, substrates are 5'- 32 P-end labeled using T4 polynucleotide kinase enzyme. Assays are performed by pre-warming a 2X concentration of purified ribozyme in ribozyme cleavage buffer (50 mM Tris-HCl, pH 7.5 at 37°C, 10 mM MgCl₂) and the cleavage reaction was initiated by adding the 2X ribozyme mix to an equal volume of substrate RNA (maximum of 1-5 nM) that was also pre-warmed in cleavage buffer. As an initial screen, assays are carried out for 1 hour at 37°C using a final concentration of either 40 nM or 1 mM ribozyme, *i.e.*, ribozyme excess. The reaction is quenched by the addition of an equal volume of 95% formamide, 20 mM

30

EDTA, 0.05% bromophenol blue and 0.05% xylene cyanol after which the sample is heated to 95°C for 2 minutes, quick chilled and loaded onto a denaturing polyacrylamide gel. Substrate RNA and the specific RNA cleavage products generated by ribozyme cleavage are visualized on an autoradiograph of the gel. The percentage of cleavage is
5 determined by Phosphor Imager® quantitation of bands representing the intact substrate and the cleavage products.

Example 5: Phospholamban

Cardiac disease leading to heart failure is the leading cause of combined morbidity
10 and mortality in the developed world. Nearly twenty million people worldwide suffer from heart failure related disease. An estimated five million Americans are afflicted with congestive heart failure (CHF), with 400,000 new cases diagnosed each year. In the US, cardiac disease associated failure results in approximately 40,000 deaths per year, and is associated with an additional 250,000 deaths (Harnish, 1999, *Drug & Market*
15 *Development*, 10, 114-119). Heart failure related disease represents a major public health issue due to an overall increase in prevalence and incidence in aging populations with a greater proportion of survivors of acute myocardial infarction (AMI) (Kannel *et al.*, 1994, *Br. Heart. J.*, 72 (suppl), 3). Heart failure related disease represents the most common reason for hospitalization of elderly patients in the US. The resulting life expectancy of
20 these patients is less than that of many common cancers, with five year survival rates for men and women at only 25% and 38% respectively, and with one year mortality rates for severe heart failure at 50% (Ho *et al.*, 1993, *Circulation*, 88, 107).

Heart disease is characterized by a progressive decrease in cardiac output resulting from insufficient pumping activity of the diseased heart. The resulting venous back-
25 pressure results in peripheral and pulmonary dysfunctional congestion. The heart responds to a variety of mechanical, hemodynamic, hormonal, and pathological stimuli by increasing muscle mass in response to an increased demand for cardiac output. The resulting transformation of heart tissue (myocardial hypertrophy) can arise as a result of genetic, physiologic, and environmental factors, and represents an early indication of
30 clinical heart disease and an important risk factor for subsequent heart failure (Hunter and Chien, 1999, *New England J. of Medicine*, 99, 313-322).

Coronary heart disease is a predominant factor in the development of the cardiac disease state, along with prior AMI, hypertension, diabetes mellitus, and valvular heart disease. Diagnosis of cardiac disease includes determination of coronary heart disease associated left ventricular systolic dysfunction (LVSD) and/or left ventricular diastolic dysfunction (LVDD) by echocardiographic imaging (Cleland, 1997, *Dis Management Health Outcomes*, 1, 169). Promising diagnosis may also rely on assaying atrial natriuretic peptide (ANP) and brain natriuretic peptide (BNP) concentrations. ANP and BNP levels are indicative of the level of ventricular dysfunction (Davidson *et al.*, 1996, *Am. J. Cardiol.*, 77, 828).

Current treatment strategies for cardiac disease associated failure are varied. Diuretics are often used to reduce pulmonary edema and dyspnea in patients with fluid overload, and are usually used in conjunction with angiotensin converting enzyme (ACE) inhibitors for vasodilation. Digoxin is another popular choice for treating cardiac disease as an inotropic agent, however, doubts remain concerning the long-term efficacy and safety of Digoxin (Harnish, 1999, *Drug & Market Development*, 10, 114-119). Carvedilol, a beta-blocker, has been introduced to complement the above treatments in order to slow down the progression of cardiac disease. Antiarrhythmic agents can be used in order to reduce the risk of sudden death in patients suffering from cardiac disease. Lastly, heart transplants have been effective in the treatment of patients with advanced stages of cardiac disease, however, the limited supply of donor hearts greatly limits the scope of this treatment to the broad population (Harnish, 1999, *Drug & Market Development*, 10, 114-119).

Whereby the above treatment strategies can all improve morbidity and mortality associated with cardiac disease, the only existing definitive approach to curing the diseased heart is replacement by transplant. Even a healthy, transplanted heart can become diseased in response to the various stresses of mechanical, hemodynamic, hormonal, and pathological stimuli associated with extrinsic risk factors. As such there exists the need for therapeutics effective in reversing the physiological changes associated with cardiac disease.

Myocardial hypertrophy and apoptosis are the underlying degenerative process associated with cardiac hypertrophy and failure. A variety of signaling pathways are involved in the progression of myocardial hypertrophy and myocardial apoptosis. Genetic

studies have been instrumental in elucidating these pathways and their involvement in cardiac disease through *in vitro* assays of cardiac muscle cells and *in vivo* studies of genetically engineered animals.

- Studies in which the expression of specific genes have been altered in cardiac myocytes have shown that specific peptide hormones, growth factors, and cytokines can activate various features of the hypertrophic response (Hunter and Chien, 1999, *New England J of Medicine*, 99, 313-322). Particular substances that have been characterized from these studies include potential therapeutic and molecular targets involved in heart failure. Hunter *et al.*, in Chien, KR, ed. *Molecular basis of heart disease: a companion to Braunwald's Heart Disease*, Philadelphia: W.B. Saunders, 1999:211-250, describe classes of therapeutic and molecular targets involved in heart failure including:
1. Endothelin 1 and angiotensin II receptor antagonists, and antagonists of ras, p38, and c-jun N-terminal kinase (JNK) for inhibition of pathologic hypertrophy.
 2. Insulin like growth factor I and growth hormone receptor stimulation for promotion of physiologic hypertrophy.
 3. beta-1-adrenergic receptor blockers for inhibition of neurohumoral over stimulation.
 4. Phospholamban and Sarcoplipin small molecule inhibitors for relief of sarcoplasmic reticulum calcium ATPase inhibition to provide enhancement of myocardial contractile and relaxation responses.
 5. Small molecule inhibitors of β -adrenergic receptor kinase to counteract the desensitization of G protein coupled receptor kinases in order to provide enhancement of myocardial contractile and relaxation responses.
 6. Enhancement of angiogenic growth factors (VEGF, FGF-5) for relief of energy deprivation in cardiac tissues.
 7. Promoters of myocyte survival including gp 130 ligands (cardiotrophin 1), and Neuregulin for the inhibition of apoptosis of myocytes.
 8. Inhibitors of apoptosis such as Caspase inhibitors for the inhibition of apoptosis of myocytes.
 9. Inhibitors of cytokines such as TNF-alpha for the inhibition of apoptosis of myocytes.
- Congestive heart failure, heart failure, dilated cardiomyopathy and pressure overload hypertrophy are nonlimiting examples of disorders and disease states that can be associated with the above classes of molecular targets.

The failure of cardiac contractile performance leading to cardiac disorders and disease, governed by impairment of cardiac excitation/contraction coupling, points to the importance of the signaling pathways involved in this process. The release and uptake of cytosolic Ca^{2+} by the sarcoplasmic reticulum plays an integral role in each cycle of cardiac contraction and excitation (Minamisawa *et al.*, 1999, *Cell*, 99, 313-322). The process of Ca^{2+} reuptake is mediated by the cardiac sarcoplasmic reticulum Ca^{2+} ATPase (SERCA2a). SERCA2a activity is regulated by phospholamban, a p52 muscle specific sarcoplasmic reticulum phosphoprotein (Koss *et al.*, 1996, *Circ. Res.*, 79, 1059-1063, and Simmerman *et al.*, 1998, *Physiol. Rev.*, 78, 921-947). In its active, unphosphorylated state, phospholamban is a potent inhibitor of SERCA2a activity. Phosphorylation of phospholamban at serine 16 by cyclic AMP-dependent protein kinase (PKA) or calmodulin kinase, results in the inhibition of phospholamban interaction with SERCA2a. This phosphorylation event is predominantly responsible for the proportional increase in the rate of Ca^{2+} uptake into the sarcoplasmic reticulum and resultant ventricular relaxation (Tada *et al.*, 1982, *Mol. Cell. Biochem.*, 46, 73-95, and Luo *et al.*, 1998, *J. Biol. Chem.*, 273, 4734-4739).

Since a proportional decrease in Ca^{2+} uptake is a hallmark feature of heart failure (Sordahl *et al.*, 1973, *Am. J. Physiol.*, 224, 497-502) and since an increase in the relative ratio of phospholamban to SERCA2a is an important determinant of sarcoplasmic reticulum dysfunction in heart failure (Hasenfuss, 1998, *Cardiovasc. Res.*, 37, 279-289), the targeting of phospholamban and related regulatory factors as therapeutic targets for heart disorders should prove valuable for cardiac indications.

Pystynen *et al.*, International PCT publication No. WO 99/00132, describe bisethers of 1-oxa, aza and thianaphthalen-2-ones as small molecule inhibitors of phospholamban for increasing coronary flow via direct dilation of the coronary arteries.

Pystynen *et al.*, International PCT publication No. WO 99/15523, describe bisethers of 1-oxa, aza and thianaphthalen-2-ones as small molecule inhibitors of phospholamban that are useful for treating heart failure.

The efficacy of the above mentioned treatment strategies is limited. Small molecule inhibition of a molecular target is often limited by toxicity, which can restrict dosing and overall efficacy.

He *et al.*, 1999, *Circulation*, 100, 974-980, describe endogenous expression of mutant phospholamban and phospholamban antisense RNA to investigate the corresponding effect on SERCA2a activity and cardiac myocyte contractility.

A more attractive approach to the treatment of heart disease would involve the use of
5 ribozymes and/or antisense constructs to modulate the expression of target molecules involved in heart failure. The use of nucleic acid molecules of the instant invention permits highly specific regulation of the molecular targets of interest, including phospholamban (PLN) (GenBank accession No. NM_002667), sarcolipin (SLN) (GenBank accession No. NM_003063), angiotensin II receptor (GenBank accession No.
10 U20860), endothelin 1 receptor (GenBank accession No. NM_001957), K-ras (GenBank accession No. NM_004985), p38 (GenBank accession No. AF092535), c-jun N-terminal kinase (GenBank accession No. NM_002750, L31951, NM_002753), growth hormone receptor (GenBank accession No. NM_000163), insulin-like growth factor I receptor (GenBank accession No. NM_000875), beta-1-adrenergic receptor (GenBank accession
15 No. NM_000024), β 1-adrenergic receptor kinase (GenBank accession No. NM_001619, NM_005160), VEGF receptor (GenBank accession No. U43368, M27281 X15997), fibroblast growth factor 5 (GenBank accession No. NM_004464), cardiotrophin I (GenBank accession No. NM_001330), neuregulin (GenBank accession No. AF009227), TNF-alpha (GenBank accession No. X02910 X02159), PI3 kinase (GenBank accession
20 No. NM_006218, NM_006219, U86453, NM_002649, M61906), and AKT kinase (GenBank accession No. NM_005163, M77198).

Various methods have been developed to assay phospholamban activity *in vitro* and *in vivo*. Holt *et al.*, 1999, *J. Mol. Cell. Cardiol.*, 31, 645-656, describe a cell culture model in which thyroid hormone control of contraction and the Ca^{2+} -ATPase/phospholamban
25 complex is studied in adult rat ventricular myocytes. Slack *et al.* 1997, *J. Biol. Chem.*, 272, 18862-18868, describe studies in which the ectopic expression of phospholamban in mouse fast-twitch skeletal muscle cells alters sarcoplasmic reticulum Ca^{2+} transport and muscle relaxation. MacLennan *et al.*, 1996, *Soc. Gen. Physiol. Ser.*, 51, 89-103, in a review of regulatory interactions between calcium ATPases and phospholamban describe
30 phospholamban/ Ca^{2+} -ATPase interactions in protein expressed in heterologous cell culture experiments. Cornwell *et al.*, 1991, *Mol. Pharmacol.*, 40, 923-931, describe the

regulation of sarcoplasmic reticulum protein phosphorylation by localized cyclic GMP-dependent protein kinase in vascular smooth muscle cells.

- Minamisawa *et al.*, 1999, *Cell*, 99, 313-322, describe a phospholamban knockout mouse model which affords protection from induced dilated cardiomyopathy. Dillmann *et al.*, 1999, *Am. J. Cardiol.*, 83, 89H-91H, describe a transgenic rat model for the study of altered expression of calcium regulatory proteins, including phospholamban, and their effect on myocyte contractile response. LekanneDeprez *et al.*, 1998, *J. Mol. Cell. Cardiol.*, 30, 1877-1888, describe a rat pressure-overload model to investigate alterations in gene expression of phospholamban, atrial natriuretic peptide (ANP), sarcoplasmic endoplasmic reticular calcium ATPase 2 (SERCA2), collagen III α 1, and calsequestrin (CSQ). Jones *et al.*, 1998, *J. Clin. Invest.*, 101, 1385-1393, describe a mouse model for investigating the regulation of calcium signaling in transgenic mouse cardiac myocytes overexpressing calsequestrin. In this study, the upregulation and downregulation of calcium uptake and release proteins were determined, including phospholamban. Lorenz *et al.*, 1997, *Am J. Physiol.*, 273, 6, describe a mouse model for the study of regulatory effects of phospholamban on cardiac function in intact mice. This study makes use of animal models with altered levels of phospholamban to permit *in vivo* evaluation of the physiological role of phospholamban. Arai *et al.*, 1996, *Saishin Igaku*, 51, 1095-1104, presents a review article of gene targeted animal models expressing cardiovascular abnormalities. The study of phospholamban and other protein expression modification effects in mice is presented. Wankerl *et al.*, 1995, *J. Mol. Med.*, 73, 487-496, presents a review article describing the study of calcium transport proteins in the nonfailing and failing heart. Animal models investigating the major calcium handling myocardial proteins, including phospholamban, are described. These models, as well as others, may be used to evaluate the effect of treatment with nucleic acid molecules of the instant invention on cardiac function. Endpoints may be, but are not limited to, left ventricular pressure, left ventricular pressure as a function of time (LVdP/dt), and mean arterial blood pressure. Endpoints will be evaluated under basal and stimulated (cardiac load) conditions.
- Particular degenerative and disease states that can be associated with phospholamban expression modulation include but are not limited to congestive heart failure, heart failure, dilated cardiomyopathy and pressure overload hypertrophy:

Digoxin, Bendrofluazide, Dofetilide, and Carvedilol are non-limiting examples of pharmaceutical agents that can be combined with or used in conjunction with the nucleic acid molecules (*e.g.* ribozymes and antisense molecules) of the instant invention. Those skilled in the art will recognize that other drugs such as diuretic and antihypertensive compounds and therapies can be similarly be readily combined with the nucleic acid molecules of the instant invention (*e.g.* ribozymes and antisense molecules) are hence within the scope of the instant invention.

Identification of Potential Target Sites in Human phospholamban RNA

The sequence of human phospholamban was screened for accessible sites using a computer folding algorithm. Regions of the RNA that did not form secondary folding structures and contained potential ribozyme and/or antisense binding/cleavage sites were identified. The sequences of these cleavage sites are shown in **Tables 24-30**.

Selection of Enzymatic Nucleic Acid Cleavage Sites in Human phospholamban RNA

Ribozyme target sites were chosen by analyzing sequences of Human phospholamban (Genbank sequence accession number: NM_002667) and prioritizing the sites on the basis of folding. Ribozymes were designed that could bind each target and were individually analyzed by computer folding (Christoffersen *et al.*, 1994 *J. Mol. Struct. Theochem*, 311, 273; Jaeger *et al.*, 1989, *Proc. Natl. Acad. Sci. USA*, 86, 7706) to assess whether the ribozyme sequences fold into the appropriate secondary structure. Those ribozymes with unfavorable intramolecular interactions between the binding arms and the catalytic core were eliminated from consideration. As noted below, varying binding arm lengths can be chosen to optimize activity. Generally, at least 5 bases on each arm are able to bind to, or otherwise interact with, the target RNA.

Chemical Synthesis and Purification of Ribozymes and Antisense for Efficient Cleavage and/or blocking of phospholamban RNA

Ribozymes and antisense constructs were designed to anneal to various sites in the RNA message. The binding arms of the ribozymes are complementary to the target site sequences described above, while the antisense constructs are fully complimentary to the target site sequences described above. The ribozymes and antisense constructs were

chemically synthesized. The method of synthesis used followed the procedure for normal RNA synthesis as described above and in Usman et al., (1987 J. Am. Chem. Soc., 109, 7845), Scaringe et al., (1990 Nucleic Acids Res., 18, 5433) and Wincott et al., *supra*, and made use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl
5 at the 5'-end, and phosphoramidites at the 3'-end. The average stepwise coupling yields were >98%.

Ribozymes and antisense constructs were also synthesized from DNA templates using bacteriophage T7 RNA polymerase (Milligan and Uhlenbeck, 1989, Methods Enzymol. 180, 51). Ribozymes and antisense constructs were purified by gel
0 electrophoresis using general methods or were purified by high pressure liquid chromatography (HPLC; see Wincott et al., *supra*; the totality of which is hereby incorporated herein by reference) and were resuspended in water. The sequences of the chemically synthesized ribozymes and antisense constructs used in this study are shown below in **Table 24-30**.

5

Ribozyme Cleavage of phospholamban RNA Target *in vitro*

Ribozymes targeted to the human phospholamban RNA are designed and synthesized as described above. These ribozymes can be tested for cleavage activity *in vitro*, for example using the following procedure. The target sequences and the nucleotide
0 location within the phospholamban RNA are given in **Tables 24-30**.

Cleavage Reactions: Full-length or partially full-length, internally-labeled target RNA for ribozyme cleavage assay is prepared by *in vitro* transcription in the presence of [α - 32 P] CTP, passed over a G 50 Sephadex column by spin chromatography and used as substrate RNA without further purification. Alternately, substrates are 5'- 32 P-end labeled
5 using T4 polynucleotide kinase enzyme. Assays are performed by pre-warming a 2X concentration of purified ribozyme in ribozyme cleavage buffer (50 mM Tris-HCl, pH 7.5 at 37°C, 10 mM MgCl₂) and the cleavage reaction was initiated by adding the 2X ribozyme mix to an equal volume of substrate RNA (maximum of 1-5 nM) that was also pre-warmed in cleavage buffer. As an initial screen, assays are carried out for 1 hour at
0 37°C using a final concentration of either 40 nM or 1 mM ribozyme, *i.e.*, ribozyme excess. The reaction is quenched by the addition of an equal volume of 95% formamide, 20 mM

EDTA, 0.05% bromophenol blue and 0.05% xylene cyanol after which the sample is heated to 95°C for 2 minutes, quick chilled and loaded onto a denaturing polyacrylamide gel. Substrate RNA and the specific RNA cleavage products generated by ribozyme cleavage are visualized on an autoradiograph of the gel. The percentage of cleavage is
5 determined by Phosphor Imager[®] quantitation of bands representing the intact substrate and the cleavage products.

Tissue distribution of BrdU-labeled antisense in mice

CD1 mice were injected with a single bolus (30 mg/kg) of a BrdU-labeled antisense
10 oligonucleotide or a similar molar amount of BrdU (as a control). At various time points (30 min, 2h and 6 h), mice were sacrificed and major tissues isolated and fixed. Distribution of antisense oligonucleotides was determined by probing with an anti-BrdU antibody and immunohistochemical staining. Tissue slices were probed with an anti-BrdU antibody followed by a reporter enzyme-conjugated second antibody and finally an enzyme
15 substrate. Visualization of the colored product by microscopy indicated nuclear staining, demonstrating effective distribution of antisense oligonucleotide in cardiac tissue.

Tissue distribution of BrdU-labeled ribozymes in monkey

Rhesus monkeys were dosed with BrdU-labeled ribozyme by intravenous bolus
20 injection at 0.1, 1.0, and 10 mg/kg once daily over five days. Saline injection was used in control animals. Animals were sacrificed and major tissues isolated and fixed. Tissue samples were probed with an anti-BrdU antibody followed by a reporter enzyme-conjugated second antibody and finally an enzyme substrate. Significant quantities of chemically modified ribozyme are detected in cardiac tissue following this dosing regimen.

25

Example 6: HBV

Chronic hepatitis B is caused by an enveloped virus, commonly known as the hepatitis B virus or HBV. HBV is transmitted via infected blood or other body fluids, especially saliva and semen, during delivery, sexual activity, or sharing of needles
30 contaminated by infected blood. Individuals may be "carriers" and transmit the infection to others without ever having experienced symptoms of the disease. Persons at highest

risk are those with multiple sex partners, those with a history of sexually transmitted diseases, parenteral drug users, infants born to infected mothers, "close" contacts or sexual partners of infected persons, and healthcare personnel or other service employees who have contact with blood. Transmission is also possible via tattooing, ear or body piercing, and acupuncture; the virus is also stable on razors, toothbrushes, baby bottles, eating utensils, and some hospital equipment such as respirators, scopes and instruments. There is no evidence that HBsAg positive food handlers pose a health risk in an occupational setting, nor should they be excluded from work. Hepatitis B has never been documented as being a food-borne disease. The average incubation period is 60 to 90 days, with a range of 45 to 180; the number of days appears to be related to the amount of virus to which the person was exposed. However, determining the length of incubation is difficult, since onset of symptoms is insidious. Approximately 50% of patients develop symptoms of acute hepatitis that last from 1 to 4 weeks. Two percent or less of these individuals develop fulminant hepatitis resulting in liver failure and death.

The determinants of severity include: (1) The size of the dose to which the person was exposed; (2) the person's age with younger patients experiencing a milder form of the disease; (3) the status of the immune system with those who are immunosuppressed experiencing milder cases; and (4) the presence or absence of co-infection with the Delta virus (hepatitis D), with more severe cases resulting from co-infection. In symptomatic cases, clinical signs include loss of appetite, nausea, vomiting, abdominal pain in the right upper quadrant, arthralgia, and tiredness/loss of energy. Jaundice is not experienced in all cases, however, jaundice is more likely to occur if the infection is due to transfusion or percutaneous serum transfer, and it is accompanied by mild pruritus in some patients. Bilirubin elevations are demonstrated in dark urine and clay-colored stools, and liver enlargement may occur accompanied by right upper-quadrant pain. The acute phase of the disease may be accompanied by severe depression, meningitis, Guillain-Barré syndrome, myelitis, encephalitis, agranulocytosis, and/or thrombocytopenia.

Hepatitis B is generally self-limiting and will resolve in approximately 6 months. Asymptomatic cases can be detected by serologic testing, since the presence of the virus leads to production of large amounts of HBsAg in the blood. This antigen is the first and most useful diagnostic marker for active infections. However, if HBsAg remains positive for 20 weeks or longer, the person is likely to remain positive indefinitely and is now a

carrier. While only 10% of persons over age 6 who contract HBV become carriers, 90% of infants infected during the first year of life do so.

Hepatitis B virus (HBV) infects over 300 million people worldwide (Imperial, 1999, *Gastroenterol. Hepatol.*, 14 (suppl), S1-5). In the United States approximately 1.25 million individuals are chronic carriers of HBV as evidenced by the fact that they have measurable hepatitis B virus surface antigen HBsAg in their blood. The risk of becoming a chronic HBsAg carrier is dependent upon the mode of acquisition of infection as well as the age of the individual at the time of infection. For those individuals with high levels of viral replication, chronic active hepatitis with progression to cirrhosis, liver failure and hepatocellular carcinoma (HCC) is common, and liver transplantation is the only treatment option for patients with end-stage liver disease from HBV.

The natural progression of chronic HBV infection over a 10 to 20 year period leads to cirrhosis in 20-to-50% of patients and progression of HBV infection to hepatocellular carcinoma has been well documented. There have been no studies that have determined sub-populations that are most likely to progress to cirrhosis and/or hepatocellular carcinoma, thus all patients have equal risk of progression.

It is important to note that the survival for patients diagnosed with hepatocellular carcinoma is only 0.9 to 12.8 months from initial diagnosis (Takahashi *et al.*, 1993, *American Journal of Gastroenterology*, 88, 240-243). Treatment of hepatocellular carcinoma with chemotherapeutic agents has not proven effective and only 10% of patients will benefit from surgery due to extensive tumor invasion of the liver (Trinchet *et al.*, 1994, *Presse Medicine*, 23, 831-833). Given the aggressive nature of primary hepatocellular carcinoma, the only viable treatment alternative to surgery is liver transplantation (Pichlmayr *et al.*, 1994, *Hepatology*, 20, 33S-40S).

Upon progression to cirrhosis, patients with chronic HCV infection present with clinical features, which are common to clinical cirrhosis regardless of the initial cause (D'Amico *et al.*, 1986, *Digestive Diseases and Sciences*, 31, 468-475). These clinical features may include: bleeding esophageal varices, ascites, jaundice, and encephalopathy (Zakim D, Boyer TD. *Hepatology a textbook of liver disease*, Second Edition Volume 1. 1990 W.B. Saunders Company. Philadelphia). In the early stages of cirrhosis, patients are classified as compensated, meaning that although liver tissue damage has occurred, the patient's liver is still able to detoxify metabolites in the blood-stream. In addition, most

patients with compensated liver disease are asymptomatic and the minority with symptoms report only minor symptoms such as dyspepsia and weakness. In the later stages of cirrhosis, patients are classified as decompensated meaning that their ability to detoxify metabolites in the bloodstream is diminished and it is at this stage that the clinical features described above will present.

In 1986, D'Amico *et al.* described the clinical manifestations and survival rates in 1155 patients with both alcoholic and viral associated cirrhosis (D'Amico *supra*). Of the 1155 patients, 435 (37%) had compensated disease although 70% were asymptomatic at the beginning of the study. The remaining 720 patients (63%) had decompensated liver disease with 78% presenting with a history of ascites, 31% with jaundice, 17% had bleeding and 16% had encephalopathy. Hepatocellular carcinoma was observed in six (0.5%) patients with compensated disease and in 30 (2.6%) patients with decompensated disease.

Over the course of six years, the patients with compensated cirrhosis developed clinical features of decompensated disease at a rate of 10% per year. In most cases, ascites was the first presentation of decompensation. In addition, hepatocellular carcinoma developed in 59 patients who initially presented with compensated disease by the end of the six-year study.

With respect to survival, the D'Amico study indicated that the five-year survival rate for all patients on the study was only 40%. The six-year survival rate for the patients who initially had compensated cirrhosis was 54% while the six-year survival rate for patients who initially presented with decompensated disease was only 21%. There were no significant differences in the survival rates between the patients who had alcoholic cirrhosis and the patients with viral related cirrhosis. The major causes of death for the patients in the D'Amico study were liver failure in 49%; hepatocellular carcinoma in 22%; and, bleeding in 13% (D'Amico *supra*).

Hepatitis B virus is a double-stranded circular DNA virus. It is a member of the Hepadnaviridae family. The virus consists of a central core that contains a core antigen (HBcAg) surrounded by an envelope containing a surface protein/surface antigen (HBsAg) and is 42 nm in diameter. It also contains an e antigen (HBeAg) which, along with HBcAg and HBsAg, is helpful in identifying this disease

In HBV virions, the genome is found in an incomplete double-stranded form. HBV uses a reverse transcriptase to transcribe a positive-sense full length RNA version of its genome back into DNA. This reverse transcriptase also contains DNA polymerase activity and thus begins replicating the newly synthesized minus-sense DNA strand. However, it
5 appears that the core protein encapsidates the reverse-transcriptase/polymerase before it completes replication.

From the free-floating form, the virus must first attach itself specifically to a host cell membrane. Viral attachment is one of the crucial steps which determines host and tissue specificity. However, currently there are no in vitro cell-lines that can be infected by
10 HBV. There are some cells lines, such as HepG2, which can support viral replication only upon transient or stable transfection using HBV DNA.

After attachment, fusion of the viral envelope and host membrane must occur to allow the viral core proteins containing the genome and polymerase to enter the cell. Once inside, the genome is translocated to the nucleus where it is repaired and cyclized.

5 The complete closed circular DNA genome of HBV remains in the nucleus and gives rise to four transcripts. These transcripts initiate at unique sites but share the same 3'-ends. The 3.5-kb pregenomic RNA serves as a template for reverse transcription and also encodes the nucleocapsid protein and polymerase. A subclass of this transcript with a 5'-end extension codes for the precore protein that, after processing, is secreted as HBV e
10 antigen. The 2.4-kb RNA encompasses the pre-S1 open reading frame (ORF) that encodes the large surface protein. The 2.1-kb RNA encompasses the pre-S2 and S ORFs that encode the middle and small surface proteins, respectively. The smallest transcript (~0.8-kb) codes for the X protein, a transcriptional activator.

Multiplication of the HBV genome begins within the nucleus of an infected cell.
15 RNA polymerase II transcribes the circular HBV DNA into greater-than-full length mRNA. Since the mRNA is longer than the actual complete circular DNA, redundant ends are formed. Once produced, the pregenomic RNA exits the nucleus and enters the cytoplasm.

The packaging of pregenomic RNA into core particles is triggered by the binding of
20 the HBV polymerase to the 5' epsilon stem-loop. RNA encapsidation is believed to occur as soon as binding occurs. The HBV polymerase also appears to require associated core protein in order to function. The HBV polymerase initiates reverse transcription from the

5' epsilon stem-loop three to four base pairs at which point the polymerase and attached nascent DNA are transferred to the 3' copy of the DR1 region. Once there, the (-)DNA is extended by the HBV polymerase while the RNA template is degraded by the HBV polymerase RNase H activity. When the HBV polymerase reaches the 5' end, a small stretch of RNA is left undigested by the RNase H activity. This segment of RNA is comprised of a small sequence just upstream and including the DR1 region. The RNA oligomer is then translocated and annealed to the DR2 region at the 5' end of the (-)DNA. It is used as a primer for the (+)DNA synthesis which is also generated by the HBV polymerase. It appears that the reverse transcription as well as plus strand synthesis may occur in the completed core particle.

Since the pregenomic RNA is required as a template for DNA synthesis, this RNA is an excellent target for ribozyme cleavage. Nucleoside analogues that have been documented to inhibit HBV replication target the reverse transcriptase activity needed to convert the pregenomic RNA into DNA. Ribozyme cleavage of the pregenomic RNA template would be expected to result in a similar inhibition of HBV replication. Further, targeting the 3'-end of the pregenomic RNA that is common to all HBV transcripts could result in reduction of all HBV gene products and an additional level of inhibition of HBV replication.

As previously mentioned, HBV does not infect cells in culture. However, transfection of HBV DNA (either as a head-to-tail dimer or as an "overlength" genome of >100%) into HuH7 or Hep G2 hepatocytes results in viral gene expression and production of HBV virions released into the media. Thus, HBV replication competent DNA would be co-transfected with ribozymes in cell culture. Such an approach has been used to report intracellular ribozyme activity against HBV (zu Putlitz, *et al.*, 1999, *J. Virol.*, 73, 5381-5387, and Kim *et al.*, 1999, *Biochem. Biophys. Res. Commun.*, 257, 759-765). In addition, stable hepatocyte cell lines have been generated that express HBV. In these cells only ribozyme would need to be delivered; however, a delivery screen would need to be performed. In addition, stable hepatocyte cell lines have been generated that express HBV.

Intracellular HBV gene expression can be assayed by a Taqman® assay for HBV RNA or by ELISA for HBV protein. Extracellular virus can be assayed by PCR for DNA or ELISA for protein. Antibodies are commercially available for HBV surface antigen and

core protein. A secreted alkaline phosphatase expression plasmid can be used to normalize for differences in transfection efficiency and sample recovery.

There are several small animal models to study HBV replication. One is the transplantation of HBV-infected liver tissue into irradiated mice. Viremia (as evidenced
5 by measuring HBV DNA by PCR) is first detected 8 days after transplantation and peaks between 18 – 25 days (Ilan *et al.*, 1999, *Hepatology*, 29, 553-562).

Transgenic mice that express HBV have also been used as a model to evaluate potential anti-virals. HBV DNA is detectable in both liver and serum (Morrey *et al.*, 1999, *Antiviral Res.*, 42, 97-108).

10 An additional model is to establish subcutaneous tumors in nude mice with Hep G2 cells transfected with HBV. Tumors develop in about 2 weeks after inoculation and express HBV surface and core antigens. HBV DNA and surface antigen is also detected in the circulation of tumor-bearing mice (Yao *et al.*, 1996, *J. Viral Hepat.*, 3, 19-22).

Woodchuck hepatitis virus (WHV) is closely related to HBV in its virus structure,
15 genetic organization, and mechanism of replication. As with HBV in humans, persistent WHV infection is common in natural woodchuck populations and is associated with chronic hepatitis and hepatocellular carcinoma (HCC). Experimental studies have established that WHV causes HCC in woodchucks and woodchucks chronically infected with WHV have been used as a model to test a number of anti-viral agents. For example,
20 the nucleoside analogue 3T3 was observed to cause dose dependent reduction in virus (50% reduction after two daily treatments at the highest dose) (Hurwitz *et al.*, 1998. *Antimicrob. Agents Chemother.*, 42, 2804-2809).

Current therapeutic goals of treatment are three-fold: to eliminate infectivity and transmission of HBV to others, to arrest the progression of liver disease and improve the
25 clinical prognosis, and to prevent the development of hepatocellular carcinoma (HCC).

Interferon alpha use is the most common therapy for HBV; however, recently Lamivudine (3TC) has been approved by the FDA. Interferon alpha (IFN-alpha) is one treatment for chronic hepatitis B. The standard duration of IFN-alpha therapy is 16 weeks, however, the optimal treatment length is still poorly defined. A complete response (HBV
30 DNA negative HBeAg negative) occurs in approximately 25% of patients. Several factors have been identified that predict a favorable response to therapy including: High ALT , low HBV DNA , being female, and heterosexual orientation.

There is also a risk of reactivation of the hepatitis B virus even after a successful response, this occurs in around 5% of responders and normally occurs within 1 year.

Side effects resulting from treatment with type 1 interferons can be divided into four general categories including: Influenza-like symptoms, neuropsychiatric, laboratory
5 abnormalities, and other miscellaneous side effects. Examples of influenza-like symptoms include, fatigue, fever; myalgia, malaise, appetite loss, tachycardia, rigors, headache and arthralgias. The influenza-like symptoms are usually short-lived and tend to abate after the first four weeks of dosing (Dusheiko *et al.*, 1994, *Journal of Viral Hepatitis*, 1, 3-5). Neuropsychiatric side effects include irritability, apathy, mood changes, insomnia,
10 cognitive changes, and depression. Laboratory abnormalities include the reduction of myeloid cells, including granulocytes, platelets and to a lesser extent, red blood cells. These changes in blood cell counts rarely lead to any significant clinical sequelae. In addition, increases in triglyceride concentrations and elevations in serum alanine and aspartate aminotransferase concentration have been observed. Finally, thyroid
15 abnormalities have been reported. These thyroid abnormalities are usually reversible after cessation of interferon therapy and can be controlled with appropriate medication while on therapy. Miscellaneous side effects include nausea, diarrhea, abdominal and back pain, pruritus, alopecia, and rhinorrhea. In general, most side effects will abate after 4 to 8 weeks of therapy (Dushieko *et al.*, *supra*).

20 Lamivudine (3TC) is a nucleoside analogue, which is a very potent and specific inhibitor of HBV DNA synthesis. Lamivudine has recently been approved for the treatment of chronic Hepatitis B. Unlike treatment with interferon, treatment with 3TC does not eliminate the HBV from the patient. Rather, viral replication is controlled and chronic administration results in improvements in liver histology in over 50% of patients.
25 Phase III studies with 3TC, showed that treatment for one year was associated with reduced liver inflammation and a delay in scarring of the liver. In addition, patients treated with Lamivudine (100mg per day) had a 98 percent reduction in hepatitis B DNA and a significantly higher rate of seroconversion, suggesting disease improvements after completion of therapy. However, stopping of therapy resulted in a reactivation of HBV
30 replication in most patients. In addition recent reports have documented 3TC resistance in approximately 30% of patients.

Particular degenerative and disease states that can be associated with HBV expression modulation include but are not limited to, HBV infection, hepatitis, cancer, tumorigenesis, cirrhosis, liver failure and others.

Lamivudine (3TC), L-FMAU, adefovir dipivoxil, type 1 Interferon, therapeutic
5 vaccines, steroids, and 2'-5' Oligoadenylates are non-limiting examples of pharmaceutical agents that can be combined with or used in conjunction with the nucleic acid molecules (e.g. ribozymes and antisense molecules) of the instant invention. Those skilled in the art will recognize that other drugs such as diuretic and antihypertensive compounds or other
10 therapies can similarly and readily be combined with the nucleic acid molecules of the instant invention (e.g. ribozymes and antisense molecules) and are, therefore, within the scope of the instant invention.

Current therapies for treating HBV infection, including interferon and nucleoside analogues, are only partially effective. In addition, drug resistance to nucleoside analogues is now emerging, making treatment of chronic Hepatitis B more difficult. Thus, a need
15 exists for effective treatment of this disease which utilizes antiviral inhibitors which work by mechanisms other than those currently utilized in the treatment of both acute and chronic hepatitis B infections.

Draper, US patent No. 6,017,756, describes the use of ribozymes for the inhibition of Hepatitis B Virus.

20 Passman *et al.*, 2000, *Biochem. Biophys. Res. Commun.*, 268(3), 728-733.; Gan *et al.*, 1998, *J. Med. Coll. PLA*, 13(3), 157-159.; Li *et al.*, 1999, *Jiefangjun Yixue Zazhi*, 24(2), 99-101.; Putlitz *et al.*, 1999, *J. Virol.*, 73(7), 5381-5387.; Kim *et al.*, 1999, *Biochem. Biophys. Res. Commun.*, 257(3), 759-765.; Xu *et al.*, 1998, *Bingdu Xuebao*, 14(4), 365-369.; Welch *et al.*, 1997, *Gene Ther.*, 4(7), 736-743.; Goldenberg *et al.*, 1997,
25 International PCT publication No. WO 97/08309, Wands *et al.*, 1997, *J. of Gastroenterology and Hepatology*, 12(suppl.), S354-S369.; Ruiz *et al.*, 1997, *BioTechniques*, 22(2), 338-345.; Gan *et al.*, 1996, *J. Med. Coll. PLA*, 11(3), 171-175.; Beck and Nassal, 1995, *Nucleic Acids Res.*, 23(24), 4954-62.; Goldenberg, 1995, International PCT publication No. WO 95/22600.; Xu *et al.*, 1993, *Bingdu Xuebao*, 9(4),
30 331-6.; Wang *et al.*, 1993, *Bingdu Xuebao*, 9(3), 278-80, all describe ribozymes that are targeted to cleave a specific HBV target site.

The enzymatic nucleic acid molecules of the instant invention exhibit a high degree of specificity for only the viral mRNA in infected cells. Nucleic acid molecules of the instant invention targeted to highly conserved sequence regions allow the treatment of many strains of human HBV with a single compound. No treatment presently exists which specifically attacks expression of the viral gene(s) that are responsible for transformation of hepatocytes by HBV.

The methods of this invention can be used to treat human hepatitis B virus infections, which include productive virus infection, latent or persistent virus infection, and HBV-induced hepatocyte transformation. The utility can be extended to other species of HBV which infect non-human animals where such infections are of veterinary importance.

Preferred target sites are genes required for viral replication, a non-limiting example includes genes for protein synthesis, such as the 5' most 1500 nucleotides of the HBV pregenomic mRNAs. For sequence references, see Renbao *et al.*, 1987, *Sci. Sin.*, 30, 507. This region controls the translational expression of the core protein (C), X protein (X) and DNA polymerase (P) genes and plays a role in the replication of the viral DNA by serving as a template for reverse transcriptase. Disruption of this region in the RNA results in deficient protein synthesis as well as incomplete DNA synthesis (and inhibition of transcription from the defective genomes). Target sequences 5' of the encapsidation site can result in the inclusion of the disrupted 3' RNA within the core virion structure and targeting sequences 3' of the encapsidation site can result in the reduction in protein expression from both the 3' and 5' fragments.

Alternative regions outside of the 5' most 1500 nucleotides of the pregenomic mRNA also make suitable targets of enzymatic nucleic acid mediated inhibition of HBV replication. Such targets include the mRNA regions that encode the viral S gene. Selection of particular target regions will depend upon the secondary structure of the pregenomic mRNA. Targets in the minor mRNAs can also be used, especially when folding or accessibility assays in these other RNAs reveal additional target sequences that are unavailable in the pregenomic mRNA species.

A desirable target in the pregenomic RNA is a proposed bipartite stem-loop structure in the 3'-end of the pregenomic RNA which is believed to be critical for viral replication (Kidd and Kidd-Ljunggren, 1996. *Nuc. Acid Res.* 24:3295-3302). The 5'end of the HBV

pregenomic RNA carries a *cis*-acting encapsidation signal, which has inverted repeat sequences that are thought to form a bipartite stem-loop structure. Due to a terminal redundancy in the pregenomic RNA, the putative stem-loop also occurs at the 3'-end. While it is the 5' copy which functions in polymerase binding and encapsidation, reverse transcription actually begins from the 3' stem-loop. To start reverse transcription, a 4 nt primer which is covalently attached to the polymerase is made, using a bulge in the 5' encapsidation signal as template. This primer is then shifted, by an unknown mechanism, to the DR1 primer binding site in the 3' stem-loop structure, and reverse transcription proceeds from that point. The 3' stem-loop, and especially the DR1 primer binding site, appear to be highly effective targets for ribozyme intervention.

Sequences of the pregenomic RNA are shared by the mRNAs for surface, core, polymerase, and X proteins. Due to the overlapping nature of the HBV transcripts, all share a common 3'-end. Ribozyme targeting this common 3'-end will thus cleave the pregenomic RNA as well as all of the mRNAs for surface, core, polymerase and X proteins.

In preferred embodiments, the invention features a method for the analysis of HBV proteins. This method is useful in determining the efficacy of HBV inhibitors. Specifically, the instant invention features an assay for the analysis of HBsAg proteins and secreted alkaline phosphatase (SEAP) control proteins to determine the efficacy of agents used to modulate HBV expression.

The method consists of coating a micro-titer plate with an antibody such as anti-HBsAg Mab (for example, Biostride B88-95-31ad,ay) at 0.1 to 10 µg/ml in a buffer (for example, carbonate buffer, such as Na₂CO₃ 15 mM, NaHCO₃ 35 mM, pH 9.5) at 4°C overnight. The microtiter wells are then washed with PBST or the equivalent thereof, (for example, PBS, 0.05% Tween 20) and blocked for 0.1-24 hr at 37° C with PBST, 1% BSA or the equivalent thereof. Following washing as above, the wells are dried (for example, at 37° C for 30 min). Biotinylated goat anti-HBsAg or an equivalent antibody (for example, Accurate YVS1807) is diluted (for example at 1:1000) in PBST and incubated in the wells (for example, 1 hr. at 37° C). The wells are washed with PBST (for example, 4x). A conjugate, (for example, Streptavidin/Alkaline Phosphatase Conjugate, Pierce 21324) is diluted to 10-10,000 ng/ml in PBST, and incubated in the wells (for example, 1 hr. at 37° C). After washing as above, a substrate (for example, p-nitrophenyl phosphate substrate,

Pierce 37620) is added to the wells, which are then incubated (for example, 1 hr. at 37° C). The optical density is then determined (for example, at 405 nm). SEAP levels are then assayed, for example, using the Great EscAPe® Detection Kit (Clontech K2041-1), as per the manufacturers instructions. In the above example, incubation times and reagent
5 concentrations may be varied to achieve optimum results, a non-limiting example is described in Example 6.

Comparison of this HBsAg ELISA method to a commercially available assay from World Diagnostics, Inc. 15271 NW 60th Ave, #201, Miami Lakes, FL 33014 (305) 827-3304 (Cat. No. EL10018) demonstrates an increase in sensitivity (signal:noise) of 3-20
10 fold.

Identification of Potential Target Sites in Human HBV RNA

The sequence of human HBV was screened for accessible sites using a computer-folding algorithm. Regions of the RNA that did not form secondary folding structures and
15 contained potential ribozyme and/or antisense binding/cleavage sites were identified. The sequences of these cleavage sites are shown in **Tables 36-43**.

Selection of Enzymatic Nucleic Acid Cleavage Sites in Human HBV RNA

Ribozyme target sites were chosen by analyzing sequences of Human HBV
20 (accession number: AF100308.1) and prioritizing the sites on the basis of folding. Ribozymes were designed that could bind each target and were individually analyzed by computer folding (Christoffersen *et al.*, 1994 *J. Mol. Struc. Theochem*, 311, 273; Jaeger *et al.*, 1989, *Proc. Natl. Acad. Sci. USA*, 86, 7706) to assess whether the ribozyme sequences fold into the appropriate secondary structure. Those ribozymes with unfavorable
25 intramolecular interactions between the binding arms and the catalytic core were eliminated from consideration. As noted herein, varying binding arm lengths can be chosen to optimize activity. Generally, at least 5 bases on each arm are able to bind to, or otherwise interact with, the target RNA.

Chemical Synthesis and Purification of Ribozymes and Antisense for Efficient Cleavage and/or blocking of HBV RNA

Ribozymes and antisense constructs were designed to anneal to various sites in the RNA message. The binding arms of the ribozymes are complementary to the target site sequences described above, while the antisense constructs are fully complementary to the target site sequences described above. The ribozymes and antisense constructs were chemically synthesized. The method of synthesis used followed the procedure for normal RNA synthesis as described above and in Usman et al., (1987 *J. Am. Chem. Soc.*, 109, 7845), Scaringe et al., (1990 *Nucleic Acids Res.*, 18, 5433) and Wincott et al., *supra*, and made use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. The average stepwise coupling yields were typically >98%.

Ribozymes and antisense constructs were also synthesized from DNA templates using bacteriophage T7 RNA polymerase (Milligan and Uhlenbeck, 1989, *Methods Enzymol.* 180, 51). Ribozymes and antisense constructs were purified by gel electrophoresis using general methods or were purified by high pressure liquid chromatography (HPLC; see Wincott et al., *supra*; the totality of which is hereby incorporated herein by reference) and were resuspended in water. The sequences of the chemically synthesized ribozymes used in this study are shown below in Table 43.

10

Ribozyme Cleavage of HBV RNA Target *in vitro*

Ribozymes targeted to the human HBV RNA are designed and synthesized as described above. These ribozymes can be tested for cleavage activity *in vitro*, for example using the following procedure. The target sequences and the nucleotide location within the HBV RNA are given in Tables 36-43.

15

Cleavage Reactions: Full-length or partially full-length, internally-labeled target RNA for ribozyme cleavage assay is prepared by *in vitro* transcription in the presence of [α -³²P] CTP, passed over a G 50 Sephadex® column by spin chromatography and used as substrate RNA without further purification. Alternately, substrates are 5'-³²P-end labeled using T4 polynucleotide kinase enzyme. Assays are performed by pre-warming a 2X concentration of purified ribozyme in ribozyme cleavage buffer (50 mM Tris-HCl, pH 7.5

20

- at 37°C, 10 mM MgCl₂) and the cleavage reaction was initiated by adding the 2X ribozyme mix to an equal volume of substrate RNA (maximum of 1-5 nM) that was also pre-warmed in cleavage buffer. As an initial screen, assays are carried out for 1 hour at 37°C using a final concentration of either 40 nM or 1 mM ribozyme, *i.e.*, ribozyme excess.
- 5 The reaction is quenched by the addition of an equal volume of 95% formamide, 20 mM EDTA, 0.05% bromophenol blue and 0.05% xylene cyanol after which the sample is heated to 95°C for 2 minutes, quick chilled and loaded onto a denaturing polyacrylamide gel. Substrate RNA and the specific RNA cleavage products generated by ribozyme cleavage are visualized on an autoradiograph of the gel. The percentage of cleavage is
- 0 determined by Phosphor Imager[®] quantitation of bands representing the intact substrate and the cleavage products.

Transfection of HepG2 Cells with psHBV-1 and Ribozymes

- The human hepatocellular carcinoma cell line Hep G2 was grown in Dulbecco's modified Eagle media supplemented with 10% fetal calf serum, 2 mM glutamine, 0.1 mM nonessential amino acids, 1 mM sodium pyruvate, 25 mM Hepes, 100 units penicillin, and 100 µg/ml streptomycin. To generate a replication competent cDNA, prior to transfection the HBV genomic sequences are excised from the bacterial plasmid sequence contained in the psHBV-1 vector (Those skilled in the art understand that other methods may be used
- 5 to generate a replication competent cDNA). This was done with an EcoRI and Hind III restriction digest. Following completion of the digest, a ligation was performed under dilute conditions (20 µg/ml) to favor intermolecular ligation. The total ligation mixture was then concentrated using Qiagen spin columns.

- Secreted alkaline phosphatase (SEAP) was used to normalize the HBsAg levels to control for transfection variability. The pSEAP2-TK control vector was constructed by ligating a Bgl II-Hind III fragment of the pRL-TK vector (Promega), containing the herpes simplex virus thymidine kinase promoter region, into Bgl II/Hind III digested pSEAP2-Basic (Clontech). Hep G2 cells were plated (3 x 10⁴ cells/well) in 96-well microtiter plates and incubated overnight. A lipid/DNA/ribozyme complex was formed containing (at final
- 0 concentrations) cationic lipid (15 µg/ml), prepared psHBV-1 (4.5 µg/ml), pSEAP2-TK (0.5 µg/ml), and ribozyme (100 µM). Following a 15 min. incubation at 37° C, the

complexes were added to the plated Hep G2 cells. Media was removed from the cells 96 hr. post-transfection for HBsAg and SEAP analysis.

Transfection of the human hepatocellular carcinoma cell line, Hep G2, with replication competent HBV DNA results in the expression of HBV proteins and the production of virions. To investigate the potential use of ribozymes for the treatment of chronic HBV infection, a series of ribozymes that target the 3' terminus of the HBV genome have been synthesized. Ribozymes targeting this region have the potential to cleave all four major HBV RNA transcripts as well as the potential to block the production of HBV DNA by cleavage of the pregenomic RNA. To test the efficacy of these HBV ribozymes, they were co-transfected with HBV genomic DNA into Hep G2 cells, and the subsequent levels of secreted HBV surface antigen (HBsAg) were analyzed by ELISA. To control for variability in transfection efficiency, a control vector which expresses secreted alkaline phosphatase (SEAP), was also co-transfected. The efficacy of the HBV ribozymes was determined by comparing the ratio of HBsAg:SEAP and/or HBeAg:SEAP to that of a scrambled attenuated control (SAC) ribozyme. Twenty-five ribozymes (RPI18341, RPI18356, RPI18363, RPI18364, RPI18365, RPI18366, RPI18367, RPI18368, RPI18369, RPI18370, RPI18371, RPI18372, RPI18373, RPI18374, RPI18303, RPI18405, RPI18406, RPI18407, RPI18408, RPI18409, RPI18410, RPI18411, RPI18418, RPI18419, and RPI18422) have been identified which cause a reduction in the levels of HBsAg and/or HBeAg as compared to the corresponding SAC ribozyme.

Example 6: Analysis of HBsAg and SEAP Levels Following Ribozyme Treatment

Immulon 4 (Dynax) microtiter wells were coated overnight at 4° C with anti-HBsAg Mab (Biostride B88-95-31ad,ay) at 1 µg/ml in Carbonate Buffer (Na₂CO₃ 15 mM, NaHCO₃ 35 mM, pH 9.5). The wells were then washed 4x with PBST (PBS, 0.05% Tween® 20) and blocked for 1 hr at 37° C with PBST, 1% BSA. Following washing as above, the wells were dried at 37° C for 30 min. Biotinylated goat ant-HBsAg (Accurate YVS1807) was diluted 1:1000 in PBST and incubated in the wells for 1 hr. at 37° C. The wells were washed 4x with PBST. Streptavidin/Alkaline Phosphatase Conjugate (Pierce 21324) was diluted to 250 ng/ml in PBST, and incubated in the wells for 1 hr. at 37° C. After washing as above, p-nitrophenyl phosphate substrate (Pierce 37620) was added to the wells, which were then incubated for 1 hr. at 37° C. The optical density at 405 nm was

then determined. SEAP levels were assayed using the Great EscApe® Detection Kit (Clontech K2041-1), as per the manufacturers instructions.

Example 7: X-gene Reporter Assay

5 The effect of ribozyme treatment on the level of transactivation of a SV40 promoter driven firefly luciferase gene by the HBV X-protein was analyzed in transfected Hep G2 cells. As a control for variability in transfection efficiency, a Renilla luciferase reporter driven by the TK promoter, which is not transactivated by the X protein, was used. Hep G2 cells were plated (3×10^4 cells/well) in 96-well microtiter plates and incubated
10 overnight. A lipid/DNA/ribozyme complex was formed containing (at final concentrations) cationic lipid (2.4 µg/ml), the X-gene vector pSBDR(2.5 µg/ml), the firefly reporter pSV40HCVluc (0.5 µg/ml), the Renilla luciferase control vector pRL-TK (0.5 µg/ml), and ribozyme (100 µM). Following a 15 min. incubation at 37° C, the complexes were added to the plated Hep G2 cells. Levels of firefly and Renilla luciferase were
15 analyzed 48 hr. post transfection, using Promega's Dual-Luciferase Assay System.

 The HBV X protein is a transactivator of a number of viral and cellular genes. Ribozymes which target the X region were tested for their ability to cause a reduction in X protein transactivation of a firefly luciferase gene driven by the SV40 promoter in
20 transfected Hep G2 cells. As a control for transfection variability, a vector containing the Renilla luciferase gene driven by the TK promoter, which is not activated by the X protein, was included in the co-transfections. The efficacy of the HBV ribozymes was determined by comparing the ratio of firefly luciferase: Renilla luciferase to that of a scrambled
25 attenuated control (SAC) ribozyme. Eleven ribozymes (RPI18365, RPI18367, RPI18368, RPI18371, RPI18372, RPI18373, RPI18405, RPI18406, RPI18411, RPI18418, RPI18423) were identified which cause a reduction in the level of transactivation of a reporter gene by the X protein, as compared to the corresponding SAC ribozyme.

Example 8: HBV transgenic mouse study

 A transgenic mouse strain (founder strain 1.3.32 with a C57B1/6 background) that
30 expresses HBV RNA and forms HBV viremia (Morrey *et al.*, 1999, *Antiviral Res.*, 42, 97-108; Guidotti *et al.*, 1995, *J. Virology*, 69, 10, 6158-6169) was utilized to study the *in vivo*

activity of ribozymes of the instant invention. This model is predictive in screening for anti-HBV agents. Ribozyme or the equivalent volume of saline was administered via a continuous s.c. infusion using Alzet® mini-osmotic pumps for 14 days. Alzet® pumps were filled with test material(s) in a sterile fashion according to the manufacturer's instructions. Prior to *in vivo* implantation, pumps were incubated at 37°C overnight (≥ 18 hours) to prime the flow modulators. On the day of surgery, animals were lightly anesthetized with a ketamine/xylazine cocktail (94 mg/kg and 6 mg/kg, respectively; 0.3 ml, IP). Baseline blood samples (200 μ l) were obtained from each animal *via* a retro-orbital bleed. A 2 cm area near the base of the tail was shaved and cleansed with betadine surgical scrub and sequentially with 70% alcohol. A 1 cm incision in the skin was made with a #15 scalpel blade or a blunt pair of scissors near the base of the tail. Forceps were used to open a pocket rostrally (*i.e.*, towards the head) by spreading apart the subcutaneous connective tissue. The pump was inserted with the delivery portal pointing away from the incision. Wounds were closed with sterile 9-mm stainless steel clips or with sterile 4-0 suture. Animals were then allowed to recover from anesthesia on a warm heating pad before being returned to their cage. Wounds were checked daily. Clips or sutures were replaced as needed. Incisions typically healed completely within 7 days post-op. Animals were then deeply anesthetized with the ketamine/xylazine cocktail (150 mg/kg and 10 mg/kg, respectively; 0.5 ml, IP) on day 14 post pump implantation. A midline thoracotomy/ laparotomy was performed to expose the abdominal cavity and the thoracic cavity. The left ventricle was cannulated at the base and animals exsanguinated using a 23G needle and 1 ml syringe. Serum was separated, frozen and analyzed for HBV DNA and antigen levels. Experimental groups were compared to the saline control group in respect to percent change from day 0 to day 14. HBV DNA was assayed by quantitative PCR

Results

Table 44 is a summary of the group designation and dosage levels used in the HBV transgenic mouse study. Baseline blood samples were obtained *via* a retroorbital bleed and animals (N=10/group) received anti-HBV ribozymes (100 mg/kg/day) as a continuous SC infusion. After 14 days, animals treated with a ribozyme targeting site 273 (RPI.18341) of

the HBV RNA showed a significant reduction in serum HBV DNA concentration, compared to the saline treated animals as measured by a quantitative PCR assay. More specifically, the saline treated animals had a 69% increase in serum HBV DNA concentrations over this 2-week period while treatment with the 273 ribozyme (RPI.18341) resulted in a 60% decrease in serum HBV DNA concentrations. Ribozymes directed against sites 1833 (RPI.18371), 1873 (RPI.18418), and 1874 (RPI.18372) decreased serum HBV DNA concentrations by 49%, 15% and 16%, respectively.

Example 7: Activity of NCH Ribozyme to inhibit HER2 gene expression

HER2 (also known as neu, erbB2 and c-erbB2) is an oncogene that encodes a 185-kDa transmembrane tyrosine kinase receptor. HER2 is a member of the epidermal growth factor receptor (EGFR) family and shares partial homology with other family members. In normal adult tissues HER2 expression is low. However, HER2 is overexpressed in at least 25-30% of breast (McGuire & Greene, 1989) and ovarian cancers (Berchuck, *et al.*, 1990). Furthermore, overexpression of HER2 in malignant breast tumors has been correlated with increased metastasis, chemoresistance and poor survival rates (Slamon *et al.*, 1987 *Science* 235: 177-182). Because HER2 expression is high in aggressive human breast and ovarian cancers, but low in normal adult tissues, it is an attractive target for ribozyme-mediated therapy (Thompson *et al.*, *supra*).

The greatest HER2 specific effects have been observed in cancer cell lines that express high levels of HER2 protein (as measured by ELISA). Specifically, in one study that treated five human breast cancer cell lines with the HER2 antibody (anti-erbB2-sFv), the greatest inhibition of cell growth was seen in three cell lines (MDA-MB-361, SKBR-3 and BT-474) that express high levels of HER2 protein. No inhibition of cell growth was observed in two cell lines (MDA-MB-231 and MCF-7) that express low levels of HER2 protein (Wright *et al.*, 1997). Another group successfully used SKBR-3 cells to show HER2 antisense oligonucleotide-mediated inhibition of HER2 protein expression and HER2 RNA knockdown (Vaughn *et al.*, 1995). Other groups have also demonstrated a decrease in the levels of HER2 protein, HER2 mRNA and/or cell proliferation in cultured cells using anti-HER2 ribozymes or antisense molecules (Suzuki, T. *et al.*, 1997; Weichen, *et al.*, 1997; Czubyko, F. *et al.*, 1997; Colomer, *et al.*, 1994; Betram *et al.*, 1994). Because cell lines that express higher levels of HER2 have been more sensitive to anti-

HER2 agents, we are pursuing several medium to high expressing cell lines, including SKBR-3 and T47D, for ribozyme screens in cell culture.

A variety of endpoints have been used in cell culture models to look at HER2-mediated effects after treatment with anti-HER2 agents. Phenotypic endpoints include inhibition of cell proliferation, apoptosis assays and reduction of HER2 protein expression. Because overexpression of HER2 is directly associated with increased proliferation of breast and ovarian tumor cells, a proliferation endpoint for cell culture assays will be our primary screen. There are several methods by which this endpoint can be measured. Following treatment of cells with ribozymes, cells are allowed to grow (typically 5 days) after which either the cell viability, the incorporation of [³H] thymidine into cellular DNA and/or the cell density can be measured. The assay of cell density is very straightforward and can be done in a 96-well format using commercially available fluorescent nucleic acid stains (such as Syto 13 or CyQuant). The assay using CyQuant is in place at RPI and is currently being employed to screen ~100 ribozymes targeting HER2 (details below).

As a secondary, confirmatory endpoint a ribozyme-mediated decrease in the level of HER2 protein expression can be evaluated using a HER2-specific ELISA.

Validation of Cell Lines and Ribozyme Treatment Conditions

Two human breast cancer cell lines (T47D and SKBR-3) that are known to express medium to high levels of HER2 protein, respectively, were considered for ribozyme screening. In order to validate these cell lines for HER2-mediated sensitivity, both cell lines were treated with the HER2 specific antibody, Herceptin® (Genentech) and its effect on cell proliferation was determined. Herceptin was added to cells at concentrations ranging from 0–8 µM in medium containing either no serum (OptiMem), 0.1% or 0.5% FBS and efficacy was determined *via* cell proliferation. Maximal inhibition of proliferation (~50%) in both cell lines was observed after addition of Herceptin at 0.5 nM in medium containing 0.1% or no FBS. The fact that both cell lines are sensitive to an anti-HER2 agent (Herceptin) supports their use in experiments testing anti-HER2 ribozymes.

Prior to ribozyme screening, the choice of the optimal lipid(s) and conditions for ribozyme delivery was determined empirically for each cell line. Applicant has established a panel of proprietary lipids that can be used to deliver ribozymes to cultured cells and are

very useful for cell proliferation assays that are typically 3-5 days in length. Initially, this panel of proprietary lipid delivery vehicles was screened in SKBR-3 and T47D cells using previously established control oligonucleotides. Specific lipids and conditions for optimal delivery were selected for each cell line based on these screens. These conditions were
5 used to deliver HER2 specific ribozymes to cells for primary (inhibition of cell proliferation) and secondary (decrease in HER2 protein) efficacy endpoints.

Primary Screen: Inhibition of Cell Proliferation

Although optimal ribozyme delivery conditions were determined for two cell lines,
10 the SKBR-3 cell line were be used for the initial screen because it has the higher level of HER2 protein, and thus should be most susceptible to a HER2-specific ribozyme. Follow-up studies can be carried out in T47D cells to confirm leads as necessary.

Ribozyme screens were be performed using an automated, high throughput 96-well cell proliferation assay. Cell proliferation were measured over a 5-day treatment period
5 using the nucleic acid stain CyQuant for determining cell density. The growth of cells treated with ribozyme/lipid complexes were compared to both untreated cells and to cells treated with Scrambled-arm Attenuated core Controls (SAC; or IA; Figure 8). SACs can no longer bind to the target site due to the scrambled arm sequence and have nucleotide changes in the core that greatly diminish ribozyme cleavage. These SACs are used to
10 determine non-specific inhibition of cell growth caused by ribozyme chemistry (*i.e.* multiple 2' *O*-Me modified nucleotides, a single 2' *C*-allyl uridine, 4 phosphorothioates and a 3' inverted abasic). Lead ribozymes are chosen from the primary screen based on their ability to inhibit cell proliferation in a specific manner. Dose response assays are carried out on these leads and a subset was advanced into a secondary screen using the
15 level of HER2 protein as an endpoint.

Secondary Screen: Decrease in HER2 Protein

A secondary screen that measures the effect of anti-HER2 ribozymes on HER2 protein levels is used to support preliminary findings. A robust HER2 ELISA for both
20 T47D and SKBR-3 cells has been established and is available for use as an additional endpoint.

Ribozyme Mechanism Assays

A Taqman assay for measuring the ribozyme-mediated decrease in HER2 RNA has also been established. This assay is based on PCR technology and can measure in real time the production of HER2 mRNA relative to a standard cellular mRNA such as GAPDH. This RNA assay is used to establish proof that lead ribozymes are working through an RNA cleavage mechanism and result in a decrease in the level of HER2 mRNA, thus leading to a decrease in cell surface HER2 protein receptors and a subsequent decrease in tumor cell proliferation.

0 Animal Models

Evaluating the efficacy of anti-HER2 agents in animal models is an important prerequisite to human clinical trials. As in cell culture models, the most HER2 sensitive mouse tumor xenografts are those derived from human breast carcinoma cells that express high levels of HER2 protein. In a recent study, nude mice bearing BT-474 xenografts were sensitive to the anti-HER2 humanized monoclonal antibody Herceptin, resulting in an 80% inhibition of tumor growth at a 1 mg kg dose (ip, 2 X week for 4-5 weeks). Tumor eradication was observed in 3 of 8 mice treated in this manner (Baselga *et al.*, 1998). This same study compared the efficacy of Herceptin alone or in combination with the commonly used chemotherapeutics, paclitaxel or doxorubicin. Although, all three anti-HER2 agents caused modest inhibition of tumor growth, the greatest antitumor activity was produced by the combination of Herceptin and paclitaxel (93% inhibition of tumor growth vs 35% with paclitaxel alone). The above studies provide proof that inhibition of HER2 expression by anti-HER2 agents causes inhibition of tumor growth in animals. Lead anti-HER2 ribozymes chosen from *in vitro* assays are further tested in mouse xenograft models. Ribozymes are first tested alone and then in combination with standard chemotherapies.

Animal Model Development

Three human breast tumor cell lines (T47D, SKBR-3 and BT-474) were characterized to establish their growth curves in mice. These three cell lines have been implanted into the mammary papillae of both nude and SCID mice and primary tumor volumes are measured 3 times per week. Growth characteristics of these tumor lines using

a Matrigel implantation format will also be established. In addition, the use of two other breast cell lines that have been engineered to express high levels of HER2 are also being used. The tumor cell line(s) and implantation method that supports the most consistent and reliable tumor growth is used in animal studies testing the lead HER2 ribozyme(s).

- 5 Ribozyme are administered by daily subcutaneous injection or by continuous subcutaneous infusion from Alzet mini osmotic pumps beginning 3 days after tumor implantation and continuing for the duration of the study. Group sizes of at least 10 animals are employed. Efficacy is determined by statistical comparison of tumor volume of ribozyme-treated animals to a control group of animals treated with saline alone. Because the growth of
0 these tumors is generally slow (45-60 days), an initial endpoint will be the time in days it takes to establish an easily measurable primary tumor (i.e. 50-100 mm³) in the presence or absence of ribozyme treatment.

Clinical Summary

- 5 Breast cancer is a common cancer in women and also occurs in men to a lesser degree. The incidence of breast cancer in the United States is ~180,000 cases per year and ~46,000 die each year of the disease. In addition, 21,000 new cases of ovarian cancer per year lead to ~13,000 deaths (data from Hung *et al.*, 1995 and the Surveillance, Epidemiology and End Results Program, NCI). Ovarian cancer is a potential secondary
10 indication for anti-HER2 ribozyme therapy.

- A full review of breast cancer is given in the NCI PDQ for Breast Cancer. A brief overview is given here. Breast cancer is evaluated or "staged" on the basis of tumor size, and whether it has spread to lymph nodes and/or other parts of the body. In Stage I breast cancer, the cancer is no larger than 2 centimeters and has not spread outside of the breast.
5 In Stage II, the patient's tumor is 2-5 centimeters but cancer may have spread to the axillary lymph nodes. By Stage III, metastasis to the lymph nodes is typical, and tumors are 5 centimeters. Additional tissue involvement (skin, chest wall, ribs, muscles *etc.*) may also be noted. Once cancer has spread to additional organs of the body, it is classed as Stage IV.

- 0 Almost all breast cancers (>90%) are detected at Stage I or II, but 31% of these are already lymph node positive. The 5-year survival rate for node negative patients (with standard surgery/radiation/chemotherapy /hormone regimens) is 97%; however,

involvement of the lymph nodes reduces the 5-year survival to only 77%. Involvement of other organs (Stage III) drastically reduces the overall survival, to 22% at 5 years. Thus, chance of recovery from breast cancer is highly dependent on early detection. Because up to 10% of breast cancers are hereditary, those with a family history are considered to be at high risk for breast cancer and should be monitored very closely.

Breast cancer is highly treatable and often curable when detected in the early stages. (For a complete review of breast cancer treatments, see the NCI PDQ for Breast Cancer.) Common therapies include surgery, radiation therapy, chemotherapy and hormonal therapy. Depending upon many factors, including the tumor size, lymph node involvement and location of the lesion, surgical removal varies from lumpectomy (removal of the tumor and some surrounding tissue) to mastectomy (removal of the breast, lymph nodes and some or all of the underlying chest muscle). Even with successful surgical resection, as many as 21% of the patients may ultimately relapse (10-20 years). Thus, once local disease is controlled by surgery, adjuvant radiation treatments, chemotherapies and/or hormonal therapies are typically used to reduce the rate of recurrence and improve survival. The therapy regimen employed depends not only on the stage of the cancer at its time of removal, but other variables such the type of cancer (ductal or lobular), whether lymph nodes were involved and removed, age and general health of the patient and if other organs are involved.

Common chemotherapies include various combinations cytotoxic drugs to kill the cancer cells. These drugs include paclitaxel (Taxol), docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil *etc.* Significant toxicities are associated with these cytotoxic therapies. Well-characterized toxicities include nausea and vomiting, myelosuppression, alopecia and mucosity. Serious cardiac problems are also associated with certain of the combinations, *e.g.* doxorubin and paclitaxel, but are less common.

Testing for estrogen and progesterone receptors helps to determine whether certain anti-hormone therapies might be helpful in inhibiting tumor growth. If either or both receptors are present, therapies to interfere with the action of the hormone ligands, can be given in combination with chemotherapy and are generally continued for several years. These adjuvant therapies are called SERMs, selective estrogen receptor modulators, and they can give beneficial estrogen-like effects on bone and lipid metabolism while antagonizing estrogen in reproductive tissues. Tamoxifen is one such compound. The

primary toxic effect associated with the use of tamoxifen is a 2 to 7-fold increase in the rate of endometrial cancer. Blood clots in the legs and lung and the possibility of stroke are additional side effects. However, tamoxifen has been determined to reduce breast cancer incidence by 49% in high-risk patients and an extensive, somewhat controversial, clinical study is underway to expand the prophylactic use of tamoxifen. Another SERM, raloxifene, was also shown to reduce the incidence of breast cancer in a large clinical trial where it was being used to treat osteoporosis. In additional studies, removal of the ovaries and/or drugs to keep the ovaries from working are being tested.

Bone marrow transplantation is being studied in clinical trials for breast cancers that have become resistant to traditional chemotherapies or where >3 lymph nodes are involved. Marrow is removed from the patient prior to high-dose chemotherapy to protect it from being destroyed, and then replaced after the chemotherapy. Another type of "transplant" involves the exogenous treatment of peripheral blood stem cells with drugs to kill cancer cells prior to replacing the treated cells in the bloodstream.

One biological treatment, a humanized monoclonal anti-HER2 antibody, Herceptin (Genentech) has been approved by the FDA as an additional treatment for HER2 positive tumors. Herceptin binds with high affinity to the extracellular domain of HER2 and thus blocks its signaling action. Herceptin can be used alone or in combination with chemotherapeutics (*i.e.* paclitaxel, docetaxel, cisplatin, *etc.*) (Pegram, *et al.*, 1998). In Phase III studies, Herceptin significantly improved the response rate to chemotherapy as well as improving the time to progression (Ross & Fletcher, 1998). The most common side effects attributed to Herceptin are fever and chills, pain, asthenia, nausea, vomiting, increased cough, diarrhea, headache, dyspnea, infection, rhinitis, and insomnia. Herceptin in combination with chemotherapy (paclitaxel) can lead to cardiotoxicity (Sparano, 1999), leukopenia, anemia, diarrhea, abdominal pain and infection.

HER2 Protein Levels for Patient Screening and as a Potential Endpoint

Because elevated HER2 levels can be detected in at least 30% of breast cancers, breast cancer patients can be pre-screened for elevated HER2 prior to admission to initial clinical trials testing an anti-HER2 ribozyme. Initial HER2 levels can be determined (by ELISA) from tumor biopsies or resected tumor samples.

During clinical trials, it may be possible to monitor circulating HER2 protein by ELISA (Ross and Fletcher, 1998). Evaluation of serial blood/serum samples over the course of the anti-HER2 ribozyme treatment period could be useful in determining early indications of efficacy. In fact, the clinical course of Stage IV breast cancer was correlated with shed HER2 protein fragment following a dose-intensified paclitaxel monotherapy. In all responders, the HER2 serum level decreased below the detection limit (Luftner *et al.*).

Two cancer-associated antigens, CA27.29 and CA15.3, can also be measured in the serum. Both of these glycoproteins have been used as diagnostic markers for breast cancer. CA27.29 levels are higher than CA15.3 in breast cancer patients; the reverse is true in healthy individuals. Of these two markers, CA27.29 was found to better discriminate primary cancer from healthy subjects. In addition, a statistically significant and direct relationship was shown between CA27.29 and large vs small tumors and node positive vs node negative disease (Gion, *et al.*, 1999). Moreover, both cancer antigens were found to be suitable for the detection of possible metastases during follow-up (Rodriguez de Paterna *et al.*, 1999). Thus, blocking breast tumor growth may be reflected in lower CA27.29 and/or CA15.3 levels compared to a control group. FDA submissions for the use of CA27.29 and CA15.3 for monitoring metastatic breast cancer patients have been filed (reviewed in Beveridge, 1999). Fully automated methods for measurement of either of these markers are commercially available.

References

Baselga, J., Norton, L. Albanell, J., Kim, Y.M. and Mendelsohn, J. (1998) Recombinant humanized anti-HER2 antibody (Herceptin) enhances the antitumor activity of paclitaxel and doxorubicin against HER2/neu overexpressing human breast cancer xenografts. *Cancer Res.* 15: 2825-2831.

Berchuck, A. Kamel, A., Whitaker, R. *et al.* (1990) Overexpression of her-2/neu is associated with poor survival in advanced epithelial ovarian cancer. *Cancer Research* 50: 4087-4091.

Bertram, J. Killian, M., Brysch, W., Schlingensiepen, K.-H., and Kneba, M. (1994) Reduction of erbB2 gene product in mamma carcinoma cell lines by erbB2 mRNA-specific and tyrosine kinase consensus phosphorothioate antisense oligonucleotides. *Biochem. BioPhys. Res. Comm.* 200: 661-667.

Beveridge, R.A. (1999) Review of clinical studies of CA27.29 in breast cancer management. *Int. J. Biol. Markers* 14: 36-39.

Colomer, R., Lupu, R., Bacus, S.S. and Gelmann, E.P. (1994) *erbB-2* antisense oligonucleotides inhibit the proliferation of breast carcinoma cells with *erbB-2* oncogene amplification. *British J. Cancer* 70: 819-825.

Czubayko, F., Downing, S.G., Hsieh, S.S., Goldstein, D.J., Lu P.Y., Trapnell, B.C. and Wellstein, A. (1997) Adenovirus-mediated transduction of ribozymes abrogates HER-2/neu and pleiotrophin expression and inhibits tumor cell proliferation. *Gene Ther.* 4: 943-949.

Gion, M., Mione, R., Leon, A.E. and Dittadi, R. (1999) Comparison of the diagnostic accuracy of CA27.29 and CA15.3 in primary breast cancer. *Clin. Chem.* 45: 630-637.

Hung, M.-C., Matin, A., Zhang, Y., Xing, X., Sorgi, F., Huang, L. and Yu, D. (1995) HER-2/neu-targeting gene therapy - a review. *Gene* 159: 65-71.

Luftner, D., Schnabel, S. and Possinger, K. (1999) c-erbB-2 in serum of patients receiving fractionated paclitaxel chemotherapy. *Int. J. Biol. Markers* 14: 55-59.

McGuire, H.C. and Greene, M.I. (1989) The *neu* (c-erbB-2) oncogene. *Semin. Oncol.* 16: 148-155.

NCI PDQ/Treatment/Health Professionals/Breast Cancer:

http://cancernet.nci.nih.gov/clinpdq/soa/Breast_cancer_Physician.html

NCI PDQ/Treatment/Patients/Breast Cancer:

http://cancernet.nci.nih.gov/clinpdq/pif/Breast_cancer_Patient.html

Pegram, M.D., Lipton, A., Hayes, D.F., Weber, B.L., Baselga, J.M., Tripathy, D., Baly, D., Baughman, S.A., Twaddell, T., Glaspy, J.A. and Slamon, D.J. (1998) Phase II study of receptor-enhanced chemosensitivity using recombinant humanized anti-p185HER2/neu monoclonal antibody plus cisplatin in patients with HER2/neu-overexpressing metastatic breast cancer refractory to chemotherapy treatment. *J. Clin. Oncol.* 16: 2659-2671.

Rodriguez de Paterna, L., Arnaiz, F., Estenoz, J. Ortuno, B. and Lanzos E. (1999) Study of serum tumor markers CEA, CA15.3, CA27.29 as diagnostic parameters in patients with breast carcinoma. *Int. J. Biol. Markers* 10: 24-29.

Ross, J.S. and Fletcher, J.A. (1998) The HER-2/neu oncogene in breast cancer: Prognostic factor, predictive factor and target for therapy. *Oncologist* 3: 1998.

Slamon, D.J., Clark, G.M., Wong, S.G., Levin, W.J., Ullrich, A. and McGuire, W.L. (1987) Human breast cancer: correlation of relapse and survival with amplification of the
5 HER-2/neu oncogene. *Science* 235: 177-182.

Sparano, J.A. (1999) Doxorubicin/taxane combinations: Cardiac toxicity and pharmacokinetics. *Semin. Oncol.* 26: 14-19.

Surveillance, Epidemiology and End Results Program (SEER) Cancer Statistics Review: http://www.seer.ims.nci.nih.gov/Publications/CSR1973_1996/

0 Suzuki T., Curcio, L.D., Tsai, J. and Kashani-Sabet M. (1997) Anti-c-erb-B-2 Ribozyme for Breast Cancer. In *Methods in Molecular Medicine*, Vol. 11, Therapeutic Applications of Ribozymes, Human Press, Inc., Totowa, NJ.

aughn, J.P., Iglehart, J.D., Demirdji, S., Davis, P., Babiss, L.E., Caruthers, M.H., Marks, J.R. (1995) Antisense DNA downregulation of the ERBB2 oncogene measured by
5 a flow cytometric assay. *Proc Natl Acad Sci USA* 92: 8338-8342.

Weichen, K., Zimmer, C. and Dietel, M. (1997) Selection of a high activity c-erbB-2 ribozyme using a fusion gene of c-erbB-2 and the enhanced green fluorescent protein. *Cancer Gene Therapy* 5: 45-51.

Wright, M., Grim, J., Deshane, J., Kim, M., Strong, T.V., Siegel, G.P., Curiel, D.T.
0 (1997) An intracellular anti-erbB-2 single-chain antibody is specifically cytotoxic to human breast carcinoma cells overexpressing erbB-2. *Gene Therapy* 4: 317-322.

Applicant has designed, synthesized and tested several NCH ribozymes and HH ribozymes targeted against HER2 RNA (see for example **Tables 31 and 34**) in cell proliferation assays.

5 **Proliferation assay:**

The model proliferation assay used in the study can require a cell plating density of 2000 cells/well in 96-well plates and at least 2 cell doublings over a 5-day treatment period. To calculate cell density for proliferation assays, the FIPS (fluoro-imaging
0 processing system) method well in the art was used. This method allows for cell density measurements after nucleic acids are stained with CyQuant dye, and has the advantage of

accurately measuring cell densities over a very wide range 1,000-100,000 cells/well in 96-well format.

Ribozymes (50-200 nM) were delivered in the presence of cationic lipid at 2.0 µg/mL and inhibition of proliferation was determined on day 5 post-treatment. Two full
5 ribozyme screens were completed and 4 lead HH and 11 lead NCH ribozymes were chosen for further testing. Of the 15 lead Rzs chosen from primary screens, 4 NCH and 1 HH Rzs continued to inhibit cell proliferation in subsequent experiments. NCH Rzs against sites, 2001 (RPI No. 17236), 2783 (RPI No. 17249), 2939 (RPI No. 17251) or 3998 (RPI No. 17262) caused inhibition of proliferation ranging from 25-60% as compared to a
10 scrambled control Rz (IA; RPI No. 17263). Of the five lead Rzs, the most efficacious is the NCH Rz (RPI No. 17251) against site 2939 of HER2 RNA. An example of results from cell culture assay is shown in Figure 8. Referring to Figure 8, NCH ribozymes and a HH ribozyme targeted against HER2 RNA, are shown to cause significant inhibition of proliferation of cells. This shows that ribozymes, for instance the NCH ribozymes are
5 capable of inhibiting HER2 gene expression in mammalian cells.

Example 8: Activity of Class II (Zinzyme) nucleic acid catalysts to inhibit HER2 gene expression

Applicant has designed, synthesized and tested several class II (zinzyme) ribozymes
10 targeted against HER2 RNA (see, for example, Tables 58, 59, and 62) in cell proliferation RNA reduction assays.

Proliferation assay:

The model proliferation assay used in the study requires a cell-plating density of
15 2000-10000 cells/well in 96-well plates and at least 2 cell doublings over a 5-day treatment period. Cells used in proliferation studies were either human breast or ovarian cancer cells (SKBR-3 and SKOV-3 cells respectively). To calculate cell density for proliferation assays, the FIPS (fluoro-imaging processing system) method well known in the art was used. This method allows for cell density measurements after nucleic acids are stained
20 with CyQuant® dye, and has the advantage of accurately measuring cell densities over a very wide range 1,000-100,000 cells/well in 96-well format.

Ribozymes (50-200 nM) were delivered in the presence of cationic lipid at 2.0-5.0 µg/mL and inhibition of proliferation was determined on day 5 post-treatment. Two full ribozyme screens were completed resulting in the selection of 14 ribozymes. Class II (zinzyme) ribozymes against sites, 314 (RPI No. 18653), 443 (RPI No. 18680), 597 (RPI No. 18697), 659 (RPI No. 18682), 878 (RPI Nos. 18683 and 18654), 881 (RPI Nos. 18684 and 18685) 934 (RPI No. 18651), 972 (RPI No. 18656, 19292, 19727, 19728, and 19293), 1292 (RPI No. 18726), 1541 (RPI No. 18687), 2116 (RPI No. 18729), 2932 (RPI No. 18678), 2540 (RPI No. 18715), and 3504 (RPI No. 18710) caused inhibition of proliferation ranging from 25-80% as compared to a scrambled control ribozyme. An example of results from a cell culture assay is shown in **Figure 20**. Referring to **Figure 20**, Class II ribozymes targeted against HER2 RNA are shown to cause significant inhibition of proliferation of cells. This shows that ribozymes, for instance the Class II (zinzyme) ribozymes are capable of inhibiting HER2 gene expression in mammalian cells.

5 RNA assay:

RNA was harvested 24 hours post-treatment using the Qiagen RNeasy® 96 procedure. Real time RT-PCR (TaqMan® assay) was performed on purified RNA samples using separate primer/probe sets specific for either target HER2 RNA or control actin RNA (to normalize for differences due to cell plating or sample recovery). Results are shown as the average of triplicate determinations of HER2 to actin RNA levels post-treatment. **Figure 30** shows class II ribozyme (zinzyme) mediated reduction in HER2 RNA targeting site 972 vs a scrambled attenuated control.

Dose response assays:

Active ribozyme was mixed with binding arm-attenuated control (BAC) ribozyme to a final oligonucleotide concentration of either 100, 200 or 400 nM and delivered to cells in the presence of cationic lipid at 5.0 µg/mL. Mixing active and BAC in this manner maintains the lipid to ribozyme charge ratio throughout the dose response curve. HER2 RNA reduction was measured 24 hours post-treatment and inhibition of proliferation was determined on day 5 post-treatment. The dose response antiproliferation results are summarized in **Figure 31** and the dose-dependent reduction of HER2 RNA results are

summarized in **Figure 32**. **Figure 33** shows a combined dose response plot of both anti-proliferation and RNA reduction data for a class II ribozyme targeting site 972 of HER2 RNA (RPI 19293).

5 **Example 9: Compositions having RNA cleaving activity**

Hammerhead ribozymes are an example of catalytic RNA molecules which are able to recognize and cleave a given specific RNA substrate (Hutchins *et al.*, 1986, *Nucleic Acids Res.* 14:3627; Keese and Symons, in *Viroids and viroid-like pathogens* (J.J. Semanchik, publ., CRC-Press, Boca Raton, Florida, 1987, pages 1-47). The catalytic
0 center of hammerhead ribozymes is flanked by three stems and can be formed by adjacent sequence regions of the RNA or also by regions, which are separated from one another by many nucleotides. **Figure 6** shows a diagram of such a catalytically active hammerhead structure. The stems have been denoted I, II and III. The nucleotides are numbered according to the standard nomenclature for hammerhead ribozymes (Hertel *et al.*, 1992,
5 *Nucleic Acids Res.* 20:3252). In this nomenclature, bases are denoted by a number, which relates their position relative to the 5' side of the cleavage site. Furthermore, each base that is involved in a stem or loop region has an additional designation (which is denoted by a decimal point and then another number) that defines the position of that base within the stem or loop. A designation of A^{15.1} would indicate that this base is involved in a paired
10 region and that it is the first nucleotide in that stem going away from the core region. This accepted convention for describing hammerhead-derived ribozymes allows for the nucleotides involved in the core of the enzyme to always have the same number relative to all of the other nucleotides. The size of the stems involved in substrate binding or core formation can be any size and of any sequence, and the position of A⁹, for example, will
5 remain the same relative to all of the other core nucleotides. Nucleotides designated, for example, N¹² or N⁹^ represent an inserted nucleotide where the position of the caret (^) relative to the number denotes whether the insertion is before or after the indicated nucleotide. Thus, N¹² represents a nucleotide inserted before nucleotide position 12, and N⁹^ represents a nucleotide inserted after nucleotide position 9.

0 The consensus sequence of the catalytic core structure is described by Ruffner and Uhlenbeck, 1990, *Nucleic Acids Res.* 18:6025-6029. Perriman *et al.*, 1992, *Gene* 113:157-163, have meanwhile shown that this structure can also contain variations, for example,

naturally occurring nucleotide insertions such as N^9 and N^{12} . Thus, the positive strand of the satellite RNA of the tobacco ring-spot virus does not contain any of the two nucleotide insertions while the +RNA strand of the virusoid of the lucerne transient streak virus (vLTSV) contains a $N^9 = U$ insertion which can be mutated to C or G without loss of activity (Sheldon and Symons, 1989, *Nucleic Acids Res.* 17:5679-5685). Furthermore, in this special case, $N^7 = A$ and $R^{15.1} = A$. On the other hand, the minus strand of the carnation stunt associated viroid (-CarSV) is quite unusual since it contains both nucleotide insertions, that is $N^{12} = A$ and $N^9 = C$ (Hernandez *et al.*, 1992, *Nucleic Acids Res.* 20:6323-6329). In this viroid $N^7 = A$ and $R^{15.1} = A$. In addition, this special hammerhead structure exhibits a very effective self-catalytic cleavage despite the more open central stem.

Possible uses of hammerhead ribozymes include, for example, generation of RNA restriction enzymes and the specific inactivation of the expression of genes in, for example, animal, human or plant cells and prokaryotes, yeasts and plasmodia. A particular biomedical interest is based on the fact that many diseases, including many forms of tumors, are related to the overexpression of specific genes. Inactivating such genes by cleaving the associated mRNA represents a possible way to control and eventually treat such diseases. Moreover there is a great need to develop antiviral, antibacterial, and antifungal pharmaceutical agents. Ribozymes have potential as such anti-infective agents since RNA molecules vital to the survival of the organism can be selectively destroyed.

In addition to needing the correct hybridizing sequences for substrate binding, substrates for hammerhead ribozymes have been shown to strongly prefer the triplet $N^{16.2}U^{16.1}H^{17}$ (NUH) where N can be any nucleotide, U is uridine, and H is either adenosine, cytidine, or uridine (Koizumi *et al.*, 1988, *FEBS Lett.* 228, 228-230; Ruffner *et al.*, 1990, *Biochemistry* 29, 10695-10702; Perriman *et al.*, 1992, *Gene* 113, 157-163). NUH is sometimes designated as NUX. The fact that changes to this general rule for substrate specificity result in non-functional substrates implies that there are "non core compatible" structures which are formed when substrates are provided which deviate from the stated requirements. Evidence along these lines was recently reported by Uhlenbeck and co-workers (Uhlenbeck *et al.*, 1997, *Biochemistry* 36:1108-1114) when they demonstrated that the substitution of a G at position 17 caused a functionally catastrophic base pair between G^{17} and C^3 to form, both preventing the correct orientation of the

scissile bond for cleavage and the needed tertiary interactions of C³ (Murray *et al.*, 1995, *Biochem. J.* 311:487-494). The strong preference for a U at position 16.1 may exist for similar reasons. Many experiments have been done in an attempt to isolate ribozymes which are able to efficiently relieve the requirement of a U at position 16.1, however, attempts to find hammerhead type ribozymes which can cleave substrates having a base other than a U at position 16.1 have proven impossible (Perriman *et al.*, 1992, *Gene* 113, 157-163).

Efficient catalytic molecules with reduced or altered requirements in the cleavage region are highly desirable because their isolation would greatly increase the number of available target sequences that molecules of this type could cleave. For example, it would be desirable to have a ribozyme variant that could efficiently cleave substrates containing triplets other than N^{16.2}U^{16.1}H¹⁷ since this would increase the number of potential target cleavage sites.

Chemically modified oligonucleotides which contain a block of deoxyribonucleotides in the middle region of the molecule have potential as pharmaceutical agents for the specific inactivation of the expression of genes (Giles *et al.*, 1992, *Nucleic Acids Res.* 20:763-770). These oligonucleotides can form a hybrid DNA-RNA duplex in which the DNA bound RNA strand is degraded by RNase H. Such oligonucleotides are considered to promote cleavage of the RNA and so cannot be characterized as having an RNA-cleaving activity nor as cleaving an RNA molecule (the RNase H is cleaving). A significant disadvantage of these oligonucleotides for *in vivo* applications is their low specificity, since hybrid formation, and thus cleavage, can also take place at undesired positions on the RNA molecules.

Since, unmodified ribozymes are sensitive to degradation by RNases, chemically modified active substances have to be used in order to administer hammerhead ribozymes exogenously (discussed, for example, by Heidenreich *et al.*, 1994, *J. Biol. Chem.* 269:2131-2138; Kiehnopf *et al.*, 1994, *EMBO J.* 13:4645-4652; Paolella *et al.*, 1992, *EMBO J.* 11:1913-1919; and Usman *et al.*, 1994, *Nucleic Acids Symp. Ser.* 31:163-164).

Sproat *et al.*, U.S. Pat. No. 5,334,711, describe such chemically modified active substances based on synthetic catalytic oligonucleotide structures with a length of 35 to 40 nucleotides which are suitable for cleaving a nucleic acid target sequence and contain modified nucleotides that contain an optionally substituted alkyl, alkenyl or alkynyl group

with 1 - 10 carbon atoms at the 2'-O atom of the ribose. These oligonucleotides contain modified nucleotide building blocks and form a structure resembling a hammerhead structure. These oligonucleotides are able to cleave specific RNA substrates.

Usman *et al.*, U.S. Patent No. 5,891,684, describe enzymatic nucleic acid molecules
5 with one or more nucleotide base modification(s) in a substrate binding arm.

Thompson *et al.*, US Patent No. 5,599,704 describe enzymatic RNA molecules targeted against ErbB2/*neu*/Her2 RNA.

Sullivan *et al.*, US Patent No. 5,616,490 describe enzymatic RNA molecules targeted against protein kinase C (PKC) RNA.

0 Sioud, International PCT publication No. WO 99/63066 describe hammerhead ribozymes targeted against specific sites within protein kinase C alpha (PKC alpha), VEGF, and TNF alpha RNA.

Jarvis *et al.*, International PCT publication No. WO 98/505030, describe the synthesis of xylo-ribonucleosides and oligonucleotides comprising xylo modifications.

5 This invention relates to novel enzymatic nucleic acid molecules having an RNA-cleavage activity, as well as their use for cleaving RNA substrates *in vitro* and *in vivo*. The compositions contain an active center, the subunits of which are selected from nucleotides and/or nucleotide analogues, as well as flanking regions contributing to the formation of a specific hybridization with an RNA substrate. Preferred compositions form, in
10 combination with an RNA substrate, a structure resembling a hammerhead structure. The active center of the disclosed compositions is characterized by the presence of I^{15.1} which allows cleavage of RNA substrates having C^{16.1}. It is therefore an object of the present invention to provide compositions that cleave RNA, and in particular to provide RNA-cleaving oligomers which at the same time have a high stability, activity, and specificity.

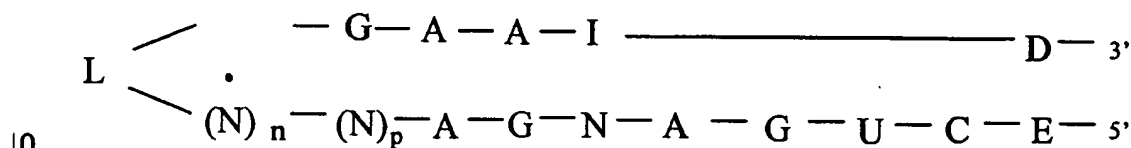
5 This invention relates to novel nucleic acid molecules with catalytic activity, which are particularly useful for cleavage of RNA or DNA or combination thereof. The nucleic acid catalysts of the instant invention are distinct from other nucleic acid catalysts known in the art. Specifically, nucleic acid catalysts of the instant invention are capable of catalyzing an intermolecular or intramolecular endonuclease reaction.

0 It is another object of the present invention to provide compositions that cleave RNA substrates having a cleavage site triplet other than N^{16.2}U^{16.1}H¹⁷ (NUH; **Figure 6**), where N is a nucleotide, U is uridine and H is adenosine, uridine or cytidine. H is used

interchangably with X. Specifically, the enzymatic nucleic acid molecule of the instant invention has an endonuclease activity to cleave RNA substrates having a cleavage triplet $N^{16.2}C^{16.1}H^{17}$ (NCH; Figure 6), where N is a nucleotide, C is cytidine and H is adenosine, uridine or cytidine. H is used interchangeably with X. In another aspect the invention

5 features an enzymatic nucleic acid molecule of the instant invention has an endonuclease activity to cleave RNA substrates having a cleavage triplet $N^{16.2}C^{16.1}N^{17}$ (NCN; Figure 6), where N is a nucleotide, C is cytidine.

In a preferred embodiment, the invention features an enzymatic nucleic acid molecule having formula 1:



where N represents independently a nucleotide or a non-nucleotide linker, which may be same or different; D and E are independently oligonucleotides of length sufficient to stably interact (*e.g.*, by forming hydrogen bonds with complementary nucleotides in the target) with a target nucleic acid molecule (the target can be an RNA, DNA or mixed

5 polymers), preferably, the length of D and E are independently between 3-20 nucleotides long, specifically, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 17, and 20; o and n are integers independently greater than or equal to 1 and preferably less than about 100, specifically 2, 3, 4, 5, 6, 7, 8, 9, 10, 15, 20, 25, 30, 35, 50, wherein if $(N)_o$ and $(N)_n$ are nucleotides, $(N)_o$ and $(N)_n$ are optionally able to interact by hydrogen bond interaction, in particular if $n=1$

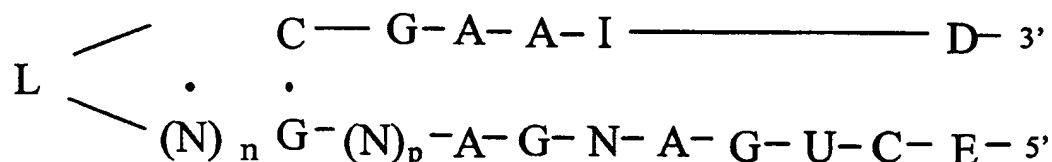
10 and $o=1$ then $(N)_n$ is preferably a purine (*e.g.*, G, and A) and $(N)_o$ is preferably a pyrimidine (*e.g.*, C and U) and $(N)_n$ preferably forms; • indicates base-paired interaction; L is a linker which may be present or absent (*i.e.*, the molecule may be assembled from two separate oligonucleotides), but when present, is a nucleotide and/or a non-nucleotide linker, which may be a single-stranded and/or double-stranded region; p is an integer 0 or

15 1, when $p=1$, $(N)_p$ is preferably A or U; and _____ represents a chemical linkage (*e.g.* a phosphate ester linkage, amide linkage, phosphorothioate linkage or others known in the art). A, U, I, C and G represent adenosine, uridine, inosine, cytidine and guanosine nucleotides, respectively. The N in 5'-CUGANGA-3' region of formula 1 is preferably U.

The nucleotides in the formula 1 are unmodified or modified at the sugar, base, and/or phosphate as known in the art.

In a preferred embodiment, the invention features an enzymatic nucleic acid molecule having formula 2:

5



0

5

0

where N represents independently a nucleotide or a non-nucleotide linker, which may be same or different; D and E are independently oligonucleotides of length sufficient to stably interact (*e.g.*, by forming hydrogen bonds with complementary nucleotides in the target) with a target nucleic acid molecule (the target can be an RNA, DNA or mixed polymers), preferably, the length of D and E are independently between 3-20 nucleotides long, specifically, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 17, and 20; o and n are integers independently greater than or equal to 0 and preferably less than about 100, specifically 2, 3, 4, 5, 6, 7, 8, 9, 10, 15, 20, 25, 30, 35, 50, wherein if (N)_o and (N)_n are nucleotides, (N)_o and (N)_n are optionally able to interact by hydrogen bond interaction; • indicates base-paired interaction; L is a linker which may be present or absent (*i.e.*, the molecule may be assembled from two separate oligonucleotides), but when present, is a nucleotide and/or a non-nucleotide linker, which may be a single-stranded and/or double-stranded region; p is an integer 0 or 1, when p=1, (N)_p is preferably A, C or U; and _____ represents a chemical linkage (*e.g.* a phosphate ester linkage, amide linkage, phosphorothioate linkage or others known in the art). A, U, I, C and G represent adenosine, uridine, inosine, cytidine and guanosine nucleotides, respectively. The N in 5'-CUGANGA-3' region of formula 2 is preferably U. The nucleotides in the formula 2 are unmodified or modified at the sugar, base, and/or phosphate as known in the art.

5

In a preferred embodiment, the I (inosine) in formula 1 and 2 is preferably a ribo-inosine or a xylo-inosine.

In yet another embodiment, the nucleotide linker (L) is a nucleic acid aptamer, such as an ATP aptamer, HIV Rev aptamer (RRE), HIV Tat aptamer (TAR) and others (for a review see Gold *et al.*, 1995, *Annu. Rev. Biochem.*, 64, 763; and Szostak & Ellington, 1993, in *The RNA World*, ed. Gesteland and Atkins, pp 511, CSH Laboratory Press). A
5 "nucleic acid aptamer" as used herein is meant to indicate nucleic acid sequence capable of interacting with a ligand. The ligand can be any natural or a synthetic molecule, including but not limited to a resin, metabolites, nucleosides, nucleotides, drugs, toxins, transition state analogs, peptides, lipids, proteins, amino acids, nucleic acid molecules, hormones, carbohydrates, receptors, cells, viruses, bacteria and others. In a preferred embodiment L
0 has the sequence 5'-GAAA-3' or 5'-GUUA-3'.

In yet another embodiment, the non-nucleotide linker (L) is as defined herein.

The term "non-nucleotide", as used herein, includes either abasic nucleotide, polyether, polyamine, polyamide, peptide, carbohydrate, lipid, or polyhydrocarbon compounds. Specific examples include those described by Seela and Kaiser, *Nucleic*
5 *Acids Res.* 1990, 18:6353 and *Nucleic Acids Res.* 1987, 15:3113; Cload and Schepartz, *J. Am. Chem. Soc.* 1991, 113:6324; Richardson and Schepartz, *J. Am. Chem. Soc.* 1991, 113:5109; Ma *et al.*, *Nucleic Acids Res.* 1993, 21:2585 and *Biochemistry* 1993, 32:1751; Durand *et al.*, *Nucleic Acids Res.* 1990, 18:6353; McCurdy *et al.*, *Nucleosides & Nucleotides* 1991, 10:287; Jschke *et al.*, *Tetrahedron Lett.* 1993, 34:301; Ono *et al.*,
10 *Biochemistry* 1991, 30:9914; Arnold *et al.*, International Publication No. WO 89/02439; Usman *et al.*, International Publication No. WO 95/06731; Dudycz *et al.*, International Publication No. WO 95/11910 and Ferentz and Verdine, *J. Am. Chem. Soc.* 1991, 113:4000, all hereby incorporated by reference herein. Non-nucleotide linkers can be any molecule, which is not an oligomeric sequence, that can be covalently coupled to an
5 oligomeric sequence. Preferred non-nucleotide linkers are oligomeric molecules formed of non-nucleotide subunits. Examples of such non-nucleotide linkers are described by Letsinger and Wu, (*J. Am. Chem. Soc.* 117:7323-7328 (1995)), Benseler *et al.*, (*J. Am. Chem. Soc.* 115:8483-8484 (1993)) and Fu *et al.*, (*J. Am. Chem. Soc.* 116:4591-4598 (1994)). Preferred non-nucleotide linkers, or subunits for non-nucleotide linkers, include
0 substituted or unsubstituted C₁-C₁₀ straight chain or branched alkyl, substituted or unsubstituted C₂-C₁₀ straight chain or branched alkenyl, substituted or unsubstituted C₂-C₁₀ straight chain or branched alkynyl, substituted or unsubstituted C₁-C₁₀ straight chain or

branched alkoxy, substituted or unsubstituted C₂-C₁₀ straight chain or branched alkenyloxy, and substituted or unsubstituted C₂-C₁₀ straight chain or branched alkynyloxy. The substituents for these preferred non-nucleotide linkers (or subunits) can be halogen, cyano, amino, carboxy, ester, ether, carboxamide, hydroxy, or mercapto. Thus, in a preferred embodiment, the invention features an enzymatic nucleic acid molecule having one or more non-nucleotide moieties, and having enzymatic activity to cleave an RNA or DNA molecule. By the term "non-nucleotide" is meant any group or compound which can be incorporated into a nucleic acid chain in the place of one or more nucleotide units, including either sugar and/or phosphate substitutions, and allows the remaining bases to exhibit their enzymatic activity. The group or compound is abasic in that it does not contain a commonly recognized nucleotide base, such as adenosine, guanine, cytosine, uracil or thymine. The terms "abasic" or "abasic nucleotide" as used herein encompass sugar moieties lacking a base or having other chemical groups in place of nucleotide base at the 1' position.

In a preferred embodiment, the invention features modified ribozymes with phosphate backbone modifications comprising one or more phosphorothioate, phosphorodithioate, methylphosphonate, morpholino, amidate carbamate, carboxymethyl, acetamidate, polyamide, sulfonate, sulfonamide, sulfamate, formacetal, thioformacetal, and/or alkylsilyl, substitutions. For a review of oligonucleotide backbone modifications see Hunziker and Leumann, 1995, *Nucleic Acid Analogues: Synthesis and Properties*, in *Modern Synthetic Methods*, VCH, 331-417, and Mesmaeker *et al.*, 1994, *Novel Backbone Replacements for Oligonucleotides*, in *Carbohydrate Modifications in Antisense Research*, ACS, 24-39.

In a further preferred embodiment of the instant invention, an inverted deoxy abasic moiety is utilized at the 3' end of the enzymatic nucleic acid molecule.

By "pyrimidines" is meant nucleotides comprising modified or unmodified derivatives of a six membered pyrimidine ring. An example of a pyrimidine is modified or unmodified uridine.

In a preferred embodiment, the nucleosides of the instant invention include, 2'-O-methyl-2,6-diaminopurine riboside; 2'-deoxy-2'-amino-2,6-diaminopurine riboside; 2'-(N-alanyl) amino-2'-deoxy-uridine; 2'-(N-phenylalanyl)amino-2'-deoxy-uridine; 2'-deoxy-2'-(N-beta-alanyl) amino ; 2'-deoxy-2'-(lysiyl) amino uridine; 2'-C-allyl uridine; 2'-O-amino-

uridine; 2'-O-methylthiomethyl adenosine; 2'-O-methylthiomethyl cytidine ; 2'-O-methylthiomethyl guanosine; 2'-O-methylthiomethyl-uridine; 2'-Deoxy-2'-(N-histidyl) amino uridine; 2'-deoxy-2'-amino-5-methyl cytidine; 2'-(N- β -carboxamidine-beta-alanyl)amino-2'-deoxy-uridine; 2'-deoxy-2'-(N-beta-alanyl)-guanosine; 2'-O-amino-adenosine; 2'-(N-lysyl)amino -2'-deoxy-cytidine; 2'-Deoxy -2'-(L-histidine) amino Cytidine; and 5-Imidazoleacetic acid 2'-deoxy-5'-triphosphate uridine.

By "oligonucleotide" as used herein is meant a molecule having two or more nucleotides. The polynucleotide can be single, double or multiple stranded and may have modified or unmodified nucleotides or non-nucleotides or various mixtures and combinations thereof.

In a preferred embodiment, the enzymatic nucleic acid molecule of formula 1 or 2 include at least three ribonucleotide residues, preferably 4, 5, 6, 7, 8, 9, and 10 ribonucleotide residues.

In preferred embodiments, the enzymatic nucleic acid of the instant invention includes one or more stretches of RNA, which provide the enzymatic activity of the molecule, linked to the non-nucleotide moiety. The necessary RNA components are known in the art (see for *e.g.*, Usman *et al.*, *supra*).

Thus, in one preferred embodiment, the invention features enzymatic nucleic acid molecules that inhibit gene expression and/or cell proliferation *in vitro* or *in vivo* (*e.g.* in patients). These chemically or enzymatically synthesized nucleic acid molecules contain substrate binding domains that bind to accessible regions of specific target nucleic acid molecules. The nucleic acid molecules also contain domains that catalyze the cleavage of target. Upon binding, the enzymatic nucleic acid molecules cleave the target molecules, preventing for example, translation and protein accumulation. In the absence of the expression of the target gene, cell proliferation, for example, is inhibited.

In another preferred embodiment, catalytic activity of the molecules described in the instant invention can be optimized as described by Draper *et al.*, *supra*. The details will not be repeated here, but include altering the length of the ribozyme binding arms, or chemically synthesizing ribozymes with modifications (base, sugar and/or phosphate) that prevent their degradation by serum ribonucleases and/or enhance their enzymatic activity (see *e.g.*, Eckstein *et al.*, International Publication No. WO 92/07065; Perrault *et al.*, 1990 *Nature* 344, 565; Pieken *et al.*, 1991 *Science* 253, 314; Usman and Cedergren, 1992

Trends in Biochem. Sci. 17, 334; Usman *et al.*, International Publication No. WO 93/15187; and Rossi *et al.*, International Publication No. WO 91/03162; Sproat, US Patent No. 5,334,711; and Burgin *et al.*, *supra*; all of these describe various chemical modifications that can be made to the base, phosphate and/or sugar moieties of enzymatic RNA molecules). Modifications which enhance their efficacy in cells, and removal of bases from stem loop structures to shorten RNA synthesis times and reduce chemical requirements are desired. (All these publications are hereby incorporated by reference herein.).

By "nucleic acid catalyst" as used herein is meant a nucleic acid molecule (*e.g.*, the molecule of formulae 1 and 2) capable of catalyzing (altering the velocity and/or rate of) a variety of reactions including the ability to repeatedly cleave other separate nucleic acid molecules (endonuclease activity) in a nucleotide base sequence-specific manner. Such a molecule with endonuclease activity may have complementarity in a substrate binding region to a specified gene target, and also has an enzymatic activity that specifically cleaves RNA or DNA in that target. That is, the nucleic acid molecule with endonuclease activity is able to intramolecularly or intermolecularly cleave RNA or DNA and thereby inactivate a target RNA or DNA molecule. This complementarity functions to allow sufficient hybridization of the enzymatic RNA molecule to the target RNA or DNA to allow the cleavage to occur. 100% complementarity is preferred, but complementarity as low as 50-75% may also be useful in this invention. The nucleic acids may be modified at the base, sugar, and/or phosphate groups. The term enzymatic nucleic acid as used herein is used interchangeably with phrases such as ribozymes, catalytic RNA, enzymatic RNA, catalytic oligonucleotides, nucleozyme, RNA enzyme, endoribonuclease, endonuclease, minizyme, oligozyme, finderone or nucleic acid catalyst. All of these terminologies describe nucleic acid molecules of the instant invention with enzymatic activity. The specific examples of enzymatic nucleic acid molecules described in the instant application are not limiting in the invention and those skilled in the art will recognize that all that is important in an enzymatic nucleic acid molecule of this invention is that it has a specific substrate binding site which is complementary to one or more of the target nucleic acid regions, and that it have nucleotide sequences within or surrounding that substrate binding site which impart a nucleic acid cleaving activity to the molecule (Cech *et al.*, U.S. Patent No. 4,987,071; Cech *et al.*, 1988, 260 *JAMA* 3030).

The enzymatic nucleic acid molecule of Formula 1 or 2 may independently comprise a cap structure which may independently be present or absent.

By "chimeric nucleic acid molecule" or "mixed polymer" is meant that, the molecule may be comprised of both modified or unmodified nucleotides.

5 In yet another preferred embodiment, the 3'-cap is selected from a group comprising, 4',5'-methylene nucleotide; 1-(beta-D-erythrofuransyl) nucleotide; 4'-thio nucleotide, carbocyclic nucleotide; 5'-amino-alkyl phosphate; 1,3-diamino-2-propyl phosphate, 3-aminopropyl phosphate; 6-aminoethyl phosphate; 1,2-aminododecyl phosphate; hydroxypropyl phosphate; 1,5-anhydrohexitol nucleotide; L-nucleotide; alpha-nucleotide; 10 modified base nucleotide; phosphorodithioate; *threo*-pentofuransyl nucleotide; acyclic 3',4'-seco nucleotide; 3,4-dihydroxybutyl nucleotide; 3,5-dihydroxypentyl nucleotide, 5'-5'-inverted nucleotide moiety; 5'-5'-inverted abasic moiety; 5'-phosphoramidate; 5'-phosphorothioate; 1,4-butanediol phosphate; 5'-amino; bridging and/or non-bridging 5'-phosphoramidate, phosphorothioate and/or phosphorodithioate, bridging or non bridging 15 methylphosphonate and 5'-mercapto moieties (for more details, see Beaucage and Iyer, 1993, *Tetrahedron* 49, 1925; incorporated by reference herein). By the term "non-nucleotide" is meant any group or compound which can be incorporated into a nucleic acid chain in the place of one or more nucleotide units, including either sugar and/or phosphate substitutions, and allows the remaining bases to exhibit their enzymatic activity. The 20 group or compound is abasic in that it does not contain a commonly recognized nucleotide base, such as adenosine, guanine, cytosine, uracil or thymine. The terms "abasic" or "abasic nucleotide" as used herein encompass sugar moieties lacking a base or having other chemical groups in place of a base at the 1' position.

In a preferred embodiment, the invention features 1-(beta-D-xylofuransyl)- 25 xypoxanthine phosphoramidite and a process for the synthesis thereof and incorporation into oligonucleotides, such as enzymatic nucleic acid molecule.

In yet another preferred embodiment, the invention features enzymatic nucleic acid molecules targeted against HER2 RNA, specifically, ribozymes in the hammerhead and NCH motifs.

30 In a preferred embodiment, the invention features enzymatic nucleic acid molecules targeted against PKC alpha RNA, specifically, ribozymes in the hammerhead and NCH motifs.

Targets, for example PKC alpha RNA, for useful ribozymes and antisense nucleic acids can be determined, for example, as described in Draper *et al.*, WO 95/04818; McSwiggen *et al.*, U.S. Patent Nos. 5,525,468 and 5,646,042, all are hereby incorporated by reference herein in their totality. Other examples include the following PCT applications, which concern inactivation of expression of disease-related genes: WO 95/23225, WO 95/13380, WO 94/02595, all incorporated by reference herein.

The specific enzymatic nucleic acid molecules described in the instant application are not limiting in the invention and those skilled in the art will recognize that all that is important in an enzymatic nucleic acid molecule of this invention is that it has a specific substrate binding site (*e.g.*, D and E of Formula 1 above) which is complementary to one or more of the target nucleic acid regions, and that it have nucleotide sequences within or surrounding that substrate binding site which impart a nucleic acid cleaving activity to the molecule.

All naturally occurring hammerhead ribozymes have an A^{15.1}-U^{16.1} base pair. In addition, it is known that substrates for ribozymes based on the consensus hammerhead sequence strongly prefer a substrate that contains an N^{16.2}U^{16.1}H¹⁷ triplet in which H¹⁷ is not a guanosine (Koizumi *et al.*, *FEBS Lett.* 228, 228-230 (1988); Ruffner *et al.*, *Biochemistry* 29, 10695-10702 (1990); Perriman *et al.*, *Gene* 113, 157-163 (1992)). Many experiments have been done in an attempt to isolate ribozymes which are able to efficiently relieve the requirement of a U at position 16.1, however, attempts to find ribozymes which can cleave substrates having a base other than a U at position 16.1 have proven largely unsuccessful (Perriman *et al.*, *Gene* 113, 157-163 1992, Singh *et al.*, *Antisense and Nucleic Acid Drug Development* 6:165-168 (1996)).

However, examination of the recently published X-ray crystal structures (Pley *et al.*, *Nature* 372:68-74 (1994), Scott *et al.*, *Cell* 81:991-1002 (1995), and Scott *et al.*, *Science* 274:2065-2069 (1996)) led to the realization that the A^{15.1}-U^{16.1} interaction is a non-standard base pair with a single hydrogen bond between the exocyclic amine (N6) of the adenosine and the 4-oxo group of the uridine. Modeling studies (based on the crystal structure) then led to the discovery that the interaction of the wild-type A^{15.1}-U^{16.1} base pair can be spatially mimicked by replacement with an I^{15.1}-C^{16.1} base pair that adopts an isostructural orientation and which preserves the required contact of the 2-keto group of C^{16.1} with A⁶ of the uridine turn. In the model, the polarity of the stabilizing hydrogen

bond between positions 15.1 and 16.1 is reversed in the $I^{15.1}-C^{16.1}$ interaction, but the correct orientation of the bases around this bond is maintained.

It has been discovered that hammerhead ribozyme analogues containing an inosine at position 15.1 readily cleave RNA substrates containing an $N^{16.2}C^{16.1}H^{17}$ triplet. Based on this, disclosed are compositions, preferably synthetic oligomers, which cleave a nucleic acid target sequence containing the triplet $N^{16.2}C^{16.1}H^{17}$. It is preferred that H^{17} is not guanosine, however, under certain circumstances, NCG triplet containing RNA can be cleaved by the ribozymes of the instant invention. The ability to cleave substrates having $N^{16.2}C^{16.1}X^{17}$ triplets effectively doubles the number of targets available for cleavage by compositions of the type disclosed.

Example 10: Synthesis of 1-(beta-D-xylofuranosyl)-xypoxanthine phosphoramidite

Referring to **Figure 9**, Inosine (1) was 5'-O-monomethoxytritylated and 2'-O-silylated under standard conditions to afford **2** (Charubala, R; Pfeleiderer, W. *Heterocycles* 1990, 30, 1141). Oxidation/reduction procedure afforded **3** in moderate yield (Matulic-Adamic, J.; Daniher, A.T.; Gonzalez, C.; Beigelman, L. *Bioorg. Med. Chem. Lett.* 1999, 9, 157): 1H NMR ($CDCl_3$) δ 12.80 (br s, 1H, NH), 8.11 (s, 1H, H-8), 8.08 (s, 1H, H-2), 7.45-6.80 (m, 14H, trityl), 5.85 (d, $J_{1',2'}=1.6$, 1H, H-1'), 4.83 (d, $J_{2',3'}=7.2$, 1H, H-2'), 4.46 (br s, 1H, 3'-OH), 4.34 (m, 1H, H-4'), 4.06 (m, 1H, H-3'), 3.77 (s, 6H, 2 x OMe), 3.60 (app d, 2H, H-5', H-5''), 0.89 (s, 9H, *t*-Bu), 0.07 (s, 3H, Me), 0.06 (s, 3H, Me).

Standard phosphitylation of **3** afforded the desired phosphoramidite **4**.

More acid stable 5'-O-MMT group is used in this particular case because applicant found that 5'-O-DMT protection is more labile in xylo nucleoside series than in ribo nucleoside series.

The xylo-inosine was incorporated into oligonucleotides using the standard procedures known in the art and as described herein.

Example 11: Activity of the xylo-Inosine-modified NCH Ribozyme

Several NCH ribozymes with xylo-inosine at position 15.1 were designed (**Figure 7**) to cleave RNA containing GCA, ACA, UCA or the CCA triplet. These ribozymes were

synthesized and purified as described herein and tested using standard RNA cleavage reaction conditions (see Table 31, for example, and see below).

The ribozymes were chemically synthesized. The method of synthesis used followed the procedure for normal RNA synthesis as described above and in Usman et al., (1987 J. Am. Chem. Soc., 109, 7845), Scaringe et al., (1990 Nucleic Acids Res., 18, 5433) and Wincott et al., supra, and made use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. The average stepwise coupling yields were >98%.

Ribozymes were purified by gel electrophoresis using general methods or were purified by high pressure liquid chromatography (HPLC; See Wincott et al., supra; the totality of which is hereby incorporated herein by reference) and were resuspended in water. The sequences of the chemically synthesized ribozymes used in this study are shown below in Table 33.

Cleavage Reactions: Full-length or partially full-length, internally-labeled target RNA for ribozyme cleavage assay is prepared by *in vitro* transcription in the presence of [α - 32 P] CTP, passed over a G 50 Sephadex column by spin chromatography and used as substrate RNA without further purification. Alternately, substrates were 5'- 32 P-end labeled using T4 polynucleotide kinase enzyme. Assays are performed by pre-warming a 2X concentration of purified ribozyme in ribozyme cleavage buffer (50 mM Tris-HCl, pH 7.5 at 37°C, 10 mM MgCl₂) and the cleavage reaction was initiated by adding the 2X ribozyme mix to an equal volume of substrate RNA (maximum of 1-5 nM) that was also pre-warmed in cleavage buffer. As an initial screen, assays are carried out for 1 hour at 37°C using a final concentration of 40 nM or 1 mM ribozyme, *i.e.*, ribozyme excess. The reaction is quenched by the addition of an equal volume of 95% formamide, 20 mM EDTA, 0.05% bromophenol blue and 0.05% xylene cyanol after which the sample is heated to 95°C for 2 minutes, quick chilled and loaded onto a denaturing polyacrylamide gel. Substrate RNA and the specific RNA cleavage products generated by ribozyme cleavage are visualized on an autoradiograph of the gel. The percentage of cleavage is determined by Phosphor Imager[®] quantitation of bands representing the intact substrate and the cleavage products.

The results of the experiments are summarized in Table 32, which shows that NCH-xylo ribozymes are catalytically active to cleave target RNA.

Example 12: Activity of NCH Ribozyme variants

5 The nucleic acid molecules of the instant invention allow for the ability to cleave a new set of 12 NCH triplets. Determination of single turnover rate constants at pH 6 of these ribozymes in the all ribo form show that with NCA type triplets, the cleavage rate is higher than at NUA sites. NCC and NUC site rates are similar, and NCU sites are slightly lower than NUU sites. Additional measurements of multiple turnover parameters of the all
0 ribo ribozymes performed under non-saturating conditions using 5nM ribozyme and changing the substrate concentration from 50 to 500 nM at pH 7.4 with 10 mM Mg⁺⁺ at 37 °C gave Km = 100 nM and kcat = 6.5 min⁻¹ for GCA vs Km = 30 nM and kcat = 2.0 min⁻¹ for GUA cleaving all ribo ribozymes. These data verify that the ribozymes with an I•C base pair are efficient catalysts in multiple turnover reactions and the relative order of
5 activity between NCH and NUH cleavers established at pH 6 (Ludwig *et al.*, 1998, *Nucleic Acids Res.*, 26, 2279-2285) remains unchanged.

To gain more insight into the structural requirements of the 15.1- 16.1 base pair of the ribozymes of the instant invention, applicant synthesized several variants of the active I-15.1 •C-16.1 structure and tested these ribozyme analogues with their corresponding
10 substrates. The influence of several core stabilization strategies on the activity of the NCH cleaving ribozymes was also investigated.

Various nucleoside analogs were incorporated at position 15.1 of the ribozyme. Cleavage activity was tested with the complementary Fl* labeled substrates at pH 7.4 in the presence of 10 mM Mg⁺⁺ under conditions of ribozyme excess (i.e. single turnover
5 conditions). The modified oligonucleotides were synthesized by standard oligonucleotide synthesis procedures. Xanthosine was protected using O-2 ,O-4 pivaloyloxymethyl groups; N,N-dimethylguanosine with 6-O-(2-nitrophenyl)-ethyl and 6-thio-inosine with S-cyanoethyl protecting groups. The cleavage activity of the ribozymes containing the 15.1 analogs is summarized in Figure 36. For comparison Figure 37 summarizes reported
0 functional group modification studies performed at the A 15.1 residue in the A-15.1 •U-16.1 context of NUH cleaving ribozymes.

Modifications at the purine 15.1 N1 and/or C6 positions (Figure 36 A, B, C)

In the 6-thio-inosine (A) (sI) 15.1 substituted ribozyme, the original (I-15.1) position 6 O•H-N (C-16.1) bonds are replaced by weaker (sI-15.1) position 6 S•H-N (C-16.1) hydrogen bonds while all other functional groups remain unchanged. Ribozymes with an
5 adenosine (B) at position 15.1 (A-15.1) are inactive with C-16.1 substrates since the ribozyme geometry requires the [A-15.1] position 6 amino group and the [C-16.1] position 4 amino group hydrogen-bond donor functional groups to be in close proximity. Similarly, low activity is observed with I-15.1 ribozymes and U-16.1 substrates, where the [I-15.1]
10 position 6 keto and [C-16.1] position 4 keto hydrogen-bond acceptor groups are opposed (Figure 37, B). Although inosine can form stable mismatch pairs with uridine in RNA duplexes or in tRNA anticodon-mRNA interactions, these results suggest that the geometry in the I•U mismatches differ from that of the A•U (or I•C) base pair in the active NUH ribozyme. Substitution of N1-Methyl-inosine (C) in place of inosine at position 15.1 leads to complete loss of cleavage activity.

Modifications at the purine 15.1 C2 and/or N3 position (Figure 36 D, E, F)

The extremely low activity observed with the G-15.1 (D) substituted analog may be explained by the formation of a G-C Watson-Crick base pair. The replacement of the I•C pair with a G•C pair can significantly distort the geometry at the 15.1-16.1 position. G-
20 15.1 N2-alkylation (E) gives only minimal recovery of catalytic activity compared to G-15.1, suggesting that the steric problems introduced by the bulky N-methyl groups may interfere with stacking interactions. The activity of this construct is significantly less than that of iso-G-15.1 (Figure 37, E) containing ribozymes in the standard A-U context. Xanthosine 15.1 (F) contains the same functional groups as inosine at the N1 and C6 sites
25 but contains an additional hydrogen-bond donor site at position N3 along with a C2 carbonyl group. The complete lack of activity seen with this construct reinforces the importance of the purine N3 acceptor functionality in transition state formation. Similarly, 3-deaza-adenosine (Figure 37, F) containing ribozymes were also inactive. The C2 carbonyl of the 15.1 purine shows no significant negative interference in iso-guanosine
30 containing 15.1 ribozymes.

Activity of modified core variants

To complete the characterization of the I•C pair containing ribozymes, the acceptance of various core substitution patterns was tested. Short substrates containing GCH and GUH (H= non G) triplets were compared using 3 different modified ribozymes. The acceptance of the U-4 2'-O-alkyl substituent is the greatest with GCA triplets while U-4= 2'-deoxy-2'-amino uridine and U-4= ribo uridine substituted ribozymes show a similar level of activity with NCH and NUH triplets. The results of this comparison are summarized in Table 64. In addition, a ribozyme construct in which ribo inosine replaces adenosine at positions 14 and 15.1 was tested which demonstrated cleavage activity.

Apart from the A-15.1 •U-16.1 to I-15 .1 •C-16.1 change that reverses the polarity of an important H-bond in the ribozyme structure, no other functional group changes at the 15.1 purine residue seem to be compatible with the requirements of efficient catalysis. The I-15.1 and A-15.1 ribozymes are equally suitable for practical applications because there are only minor differences in the acceptance of stabilizing residues.

Example 13: Activity of NCH Ribozyme to inhibit HER2 gene expression

Applicant has designed, synthesized and tested several NCH ribozymes and HH ribozymes targeted against HER2 RNA (see, for example, Tables 31 and 34) in cell proliferation assays.

Proliferation assay: The model proliferation assay used in the study can require a cell plating density of 2000 cells/well in 96-well plates and at least 2 cell doublings over a 5-day treatment period. To calculate cell density for proliferation assays, the FIPS (fluoro-imaging processing system) method well in the art was used. This method allows for cell density measurements after nucleic acids are stained with CyQuant® dye, and has the advantage of accurately measuring cell densities over a very wide range 1,000-100,000 cells/well in 96-well format.

Ribozymes (50-200 nM) were delivered in the presence of cationic lipid at 2.0 µg/mL and inhibition of proliferation was determined on day 5 post-treatment. Two full ribozyme screens were completed and 4 lead HH and 11 lead NCH ribozymes were chosen for further testing. Of the 15 lead Rzs chosen from primary screens, 4 NCH and 1

HH Rzs continued to inhibit cell proliferation in subsequent experiments. NCH Rzs against sites, 2001 (RPI No. 17236), 2783 (RPI No. 17249), 2939 (RPI No. 17251) or 3998 (RPI No. 17262) caused inhibition of proliferation ranging from 25-60% as compared to a scrambled control Rz (IA; RPI No. 17263). Of the five lead Rzs, the most efficacious is the NCH Rz (RPI No. 17251) against site 2939 of HER2 RNA. An example of results from cell culture assay is shown in Figure 3. Referring to Figure 3, NCH ribozymes and a HH ribozyme targeted against HER2 RNA are shown to cause significant inhibition of proliferation of cells. This shows that ribozymes, for instance, the NCH ribozymes are capable of inhibiting HER2 gene expression in mammalian cells.

Example 14: Activity of NCH Ribozyme to inhibit PKC alpha gene expression

The Protein Kinase C family contains twelve currently known isozymes divided into three classes: the classic, Ca^{++} dependent ($\text{PKC}\alpha$, βI , βII , γ), the novel, non- Ca^{++} dependent ($\text{PKC}\delta$, ϵ , μ , η , θ) and the atypical ($\text{PKC}\xi$, i/λ); all of which are serine/threonine kinases. These isozymes show distinct and overlapping tissue, cellular, and subcellular distribution. They aid in the regulation of cell growth and differentiation through their response to second messenger products of lipid metabolism (Blobe, *et al.*, 1996, *Cancer Surveys*, 27, 213-248). These second messengers include diacylglycerol (DAG), inositol-triphosphate (IP3), lysophospholipids, free fatty acids, and phosphatidate which act directly or in addition to changes in the Ca^{++} concentration. A simple model for $\text{PKC}\alpha$ activation follows a two step mechanism. First, membrane association of $\text{PKC}\alpha$ is through Ca^{++} and phospholipid interactions and second, the kinase is activated by interaction with DAG. An example of a signal cascade subsequent to PKC activation is PKC's phosphorylation of c-Raf, which phosphorylates MEK, which phosphorylates MAP, which phosphorylates transcription factors such as Jun and thereby activates a mitogenic program in the nucleus. There are numerous substrates for the various PKC's, one which for $\text{PKC}\alpha$ ultimately stimulates transcription factors that activate P-glycoprotein (P-gp) causing the multi-drug resistant phenotype (MDR) (Blobe, *et al.*, 1994, *Cancer and Metastasis Reviews*, 13, 411-431).

Cell Culture Review

PKC's have been implicated in tumor promotion since the discovery that these molecules can serve as receptors for tumor-promoting phorbol esters. An increase in PKC overexpression in numerous tumor cell lines and tumor tissues has also been demonstrated. PKC overexpression has been shown to be associated with increased invasion and metastasis in mouse Lewis lung carcinoma, mouse B16 melanoma (Lee *et al.*, 1997, *Molecular Carcinogenesis*, 18, 44-53), mouse mammary adenocarcinoma, mouse fibrosarcoma, human lung carcinoma (Wang and Liu, 1998, *Acta Pharmacologica Sinica*, 19, 265-268), human bladder carcinoma, human pancreatic cancer (Denham *et al.*, 1998, *Surgery*, 124, 218-223), and human gastric cancer (Dean *et al.*, 1996, *Cancer Research*, 56, 3499-3507). Mounting evidence suggests PKC α can stimulate adhesion molecule expression and can directly act on these membrane bound species as substrates, thereby modulating cellular adhesion to the extracellular matrix and increasing metastatic potential. Furthermore, human surgical specimens have demonstrated elevated PKC in breast tumors, thyroid carcinomas and melanomas (Becker *et al.*, 1990, *Oncogene*, 5, 1133-1139).

Utz *et al.*, 1994, *Int. J. Cancer*, 57, 104-110, describe a cell proliferation assay in which small molecule inhibitors of PKC demonstrate anti-proliferative activity in CCRF-VCR 1000 and KB-8511 cells with the multidrug resistant (MDR) phenotype. PKC α is overexpressed in tumor tissues that express the MDR phenotype. This phenotype is associated with the expression of a 170 kDa broad specificity drug efflux pump, P-gp. PKC α phosphorylation of P-gp has been shown *in vitro*. In addition, PKC expression correlates with resistance to doxorubicin and high P-gp levels in human renal carcinoma and non-small cell lung carcinoma. Inhibitors of PKC partially reverse the MDR phenotype and decrease phosphorylation of P-gp (Caponigro *et al.*, 1997, *Anti-Cancer Drugs*, 8, 26-33).

Dean *et al.*, 1994, *Journal of Biological Chemistry*, 269, 16416-24, describe cell culture studies in which antisense targeting of PKC α resulted in the potent inhibition of mRNA and protein expression in human lung carcinoma (A549) cells. In this study, PKC α inhibition resulted in the reduced induction of intercellular adhesion molecule 1 (ICAM-1) mRNA by phorbol esters.

Yano *et al.*, 1999, *Endocrinology*, 140, 4622-4632, describe a cell proliferation study in which down regulation of different PKC isoforms, including PKC α , results in the inhibition of insulin like growth factor I induced vascular smooth muscle cell proliferation, migration, and gene expression.

- 5 Wang *et al.*, 1999, *Experimental Cell Research*, 250, 253-263, describe cell culture studies in which antisense inhibition of PKC α results in the reversal of the transformed phenotype in human lung carcinoma (LTEPa-2) cells. In this study, the amounts of PKC α protein and total PKC activity were decreased when compared to control cells.

- Sioud and Sorensen, 1998, *Nature Biotechnology*, 16, 556-561, describe
0 hammerhead ribozyme inhibition of PKC α in rat glioma cell lines (BT4C and BT4Cn). This study demonstrated inhibition of malignant glioma cell proliferation along with the inhibition of regulatory Bcl-x_L protein expression. Bcl-x_L is overexpressed in glioma cells and is an apoptosis inhibitor. The ribozyme mediated inhibition of cell proliferation presumably results from apoptosis induction of transformed glioma cells through
5 suppression of PKC α and Bcl-x_L (Leirdal and Sioud, 1999, *British J. of Cancer*, 80, 1558-1564).

Animal Models

- Evaluating the efficacy of anti-PKC α agents in animal models is an important prerequisite to human clinical trials. A variety of mouse xenograft models using human
0 tumor cell lines have been developed using cell lines which express high levels of PKC α protein. McGraw *et al.*, 1997, *Anti-Cancer Drug Design*, 12, 315-326, describe mouse xenograft models using human breast (MDA MB-321), prostate (Du-145), colon (Colo 205, WiDr), lung (NCI H69, H209, J460, H520, A549), bladder (T-24), and melanoma (SK-mel 1) carcinoma cells. Antisense oligonucleotides targeting PKC α administered
5 intravenously following s.c. transplanted tumor cells resulted in dose dependant decreases in tumor size when compared to controls in most cases. Similar studies using T-24 bladder carcinoma, non-small cell lung carcinoma (A549), and Colo 205 colon carcinoma mouse xenografts are described in Dean *et al.*, 1996, *Biochemical Society Transactions*, 24, 623. Sioud and Sorensen, 1998, *Nature Biotechnology*, 16, 556-561, describe a rat model
0 in which inbred syngeneic BDIX rats were inoculated subcutaneously with BT4Cn glioma cells. After approximately three weeks, rats were treated with a single injection of

ribozyme targeting PKC α resulting in inhibition of tumor growth as determined by tumor size and/or weight when compared to controls. The above studies provide proof that inhibition of PKC α expression by anti-PKC α agents causes inhibition of tumor growth in animals. Lead anti-PKC α ribozymes chosen from *in vitro* assays can be further tested in
5 mouse xenograft models. Ribozymes can be first tested alone and then in combination with standard chemotherapies.

Animal Model Development

Human lung (A549, NCI H520) tumor and breast (MDA-MB 231) cell lines can be characterized to establish their growth curves in mice. These cell lines are been implanted
10 into both nude and SCID mice and primary tumor volumes are measured 3 times per week. Growth characteristics of these tumor lines using a Matrigel implantation format can also be established. In addition, the use of other cell lines that have been engineered to express high levels of PKC α can also be used. The tumor cell line(s) and implantation method that supports the most consistent and reliable tumor growth can be used in animal studies
15 to test promising PKC α ribozyme(s). Ribozymes can be administered by daily subcutaneous injection or by continuous subcutaneous infusion from Alzet mini osmotic pumps beginning 3 days after tumor implantation and continuing for the duration of the study. Group sizes of at least 10 animals are employed. Efficacy is determined by statistical comparison of tumor volume of ribozyme-treated animals to a control group of
20 animals treated with saline alone. Because the growth of these tumors is generally slow (45-60 days), an initial endpoint will be the time in days it takes to establish an easily measurable primary tumor (i.e. 50-100 mm³) in the presence or absence of ribozyme treatment.

Clinical Summary

25 *Overview*

Ribozymes targeting PKC α have strong potential to develop into useful therapeutics directed towards numerous cancer types. Lung cancer is the leading cause of cancer deaths for both men and women in the USA. The incidence of lung cancer in the United States is ~172,000 cases per year, accounting for 14% of cancer diagnoses. Approximately 158,000
30 die each year of lung cancer, accounting for 28% of all cancer deaths. Numerous other

indications exist including cancers of the bladder, colon, breast, prostate, and ovary in addition to melanoma and glioblastoma.

McGraw *et al.*, 1997, *Anti-Cancer Drug Design*, 12, 315-326, describe a Phase I trial for ISIS 3521/CGP 64128A, a PKC alpha antisense construct. In this trial, ISIS 3521/CGP 64128A was administered as either a two-hour i.v. infusion three times per week for three consecutive weeks, or as a continuous i.v. infusion for twenty-one consecutive days. The authors report that patients demonstrated excellent tolerance to the antisense compound when administered at doses of up to 2.5 mg/kg by the two-hour i.v. infusion and at 1.5 mg/kg/day by continuous i.v. infusion. In patients receiving the two-hour i.v. infusion schedule, the post-infusion plasma concentration of the compound increased proportional to the dose, and metabolites were determined to have been cleared rapidly from plasma with a half-life of thirty to forty-five minutes. These metabolites were composed of chain-shortened oligonucleotides, consistent with exonuclease-mediated degradation. No evidence of accumulation, induction, or inhibition of metabolism was found after the administration of repetitive doses.

Therapy

Treatment options for lung cancer are determined by the type and stage of the cancer and include surgery, radiation therapy, and chemotherapy. For many localized cancers, surgery is usually the treatment of choice. Because the disease has usually spread by the time it is discovered, radiation therapy and chemotherapy are often needed in combination with surgery. Chemotherapy alone or combined with radiation has replaced surgery as the treatment of choice for small cell lung cancer; on this regimen, a large percentage of patients experience remission, which in some cases is long-lasting. The 1-year relative survival rates for lung cancer have increased from 32% in 1973 to 41% in 1994, largely due to improvements in surgical techniques. The 5-year relative survival rate for all stages combined is only 14%. The survival rate is 50% for cases detected when the disease is still localized, but only 15% of lung cancers are discovered that early.

Common chemotherapies include various combinations of cytotoxic drugs to kill the cancer cells. These drugs include paclitaxel (Taxol), docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil *etc.* Significant toxicities are associated with these cytotoxic therapies. Well-characterized toxicities include nausea and vomiting,

myelosuppression, alopecia and mucosity. Serious cardiac problems are also associated with certain of the combinations, e.g. doxorubin and paclitaxel, but are less common.

Applicant has designed several NCH ribozymes targeted against PKC α RNA (Genebank accession No NM_002737) (see, for example, Table 63). These ribozymes are used first in a proliferation assay that is used to select ribozyme leads.

Proliferation assay: The model proliferation assay useful in the study can require a cell plating density of 2000 cells/well in 96-well plates and at least 2 cell doublings over a 5-day treatment period. To calculate cell density for proliferation assays, the FIPS (fluoro-imaging processing system) method well known in the art can be used. This method allows for cell density measurements after nucleic acids are stained with CyQuant® dye, and has the advantage of accurately measuring cell densities over a very wide range 1,000-100,000 cells/well in 96-well format.

Ribozymes (50-200 nM) are delivered in the presence of cationic lipid at 2.0 μ g/mL and inhibition of proliferation is determined on day 5 post-treatment. Two full ribozyme screens are usually completed and lead ribozymes are chosen for further testing. Of the lead ribozymes chosen from primary screens, ribozymes which continue to inhibit cell proliferation in subsequent experiments are selected for PKC α RNA and protein inhibition studies.

Example 15: Nucleoside Triphosphates and their incorporation into oligonucleotides

The synthesis of nucleotide triphosphates and their incorporation into nucleic acids using polymerase enzymes has greatly assisted in the advancement of nucleic acid research. The polymerase enzyme utilizes nucleotide triphosphates as precursor molecules to assemble oligonucleotides. Each nucleotide is attached by a phosphodiester bond formed through nucleophilic attack by the 3' hydroxyl group of the oligonucleotide's last nucleotide onto the 5' triphosphate of the next nucleotide. Nucleotides are incorporated one at a time into the oligonucleotide in a 5' to 3' direction. This process allows RNA to be produced and amplified from virtually any DNA or RNA templates.

Most natural polymerase enzymes incorporate standard nucleotide triphosphates into nucleic acid. For example, a DNA polymerase incorporates dATP, dTTP, dCTP, and dGTP into DNA and an RNA polymerase generally incorporates ATP, CTP, UTP, and

GTP into RNA. There are however, certain polymerases that are capable of incorporating non-standard nucleotide triphosphates into nucleic acids (Joyce, 1997, *PNAS* 94, 1619-1622, Huang et al., *Biochemistry* 36, 8231-8242).

Before nucleosides can be incorporated into RNA transcripts using polymerase
5 enzymes they must first be converted into nucleotide triphosphates which can be recognized by these enzymes. Phosphorylation of unblocked nucleosides by treatment with POCl₃ and trialkyl phosphates was shown to yield nucleoside 5'-phosphorodichloridates (Yoshikawa et al., 1969, *Bull. Chem. Soc. (Japan)* 42, 3505). Adenosine or 2'-deoxyadenosine 5'-triphosphate was synthesized by adding an additional
10 step consisting of treatment with excess tri-n-butylammonium pyrophosphate in DMF followed by hydrolysis (Ludwig, 1981, *Acta Biochim. et Biophys. Acad. Sci. Hung.* 16, 131-133).

Non-standard nucleotide triphosphates are not readily incorporated into RNA transcripts by traditional RNA polymerases. Mutations have been introduced into RNA
15 polymerase to facilitate incorporation of deoxyribonucleotides into RNA (Sousa & Padilla, 1995, *EMBO J.* 14, 4609-4621, Bonner et al., 1992, *EMBO J.* 11, 3767-3775, Bonner et al., 1994, *J. Biol. Chem.* 42, 25120-25128, Aurup et al., 1992, *Biochemistry* 31, 9636-9641).

McGee et al., International PCT Publication No. WO 95/35102, describes the incorporation of 2'-NH₂-NTP's, 2'-F-NTP's, and 2'-deoxy-2'-benzyloxyamino UTP into
20 RNA using bacteriophage T7 polymerase.

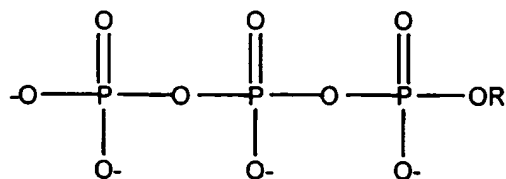
Wieczorek et al., 1994, *Bioorganic & Medicinal Chemistry Letters* 4, 987-994, describes the incorporation of 7-deaza-adenosine triphosphate into an RNA transcript using bacteriophage T7 RNA polymerase.

Lin et al., 1994, *Nucleic Acids Research* 22, 5229-5234, reports the incorporation of
25 2'-NH₂-CTP and 2'-NH₂-UTP into RNA using bacteriophage T7 RNA polymerase and polyethylene glycol containing buffer. The article describes the use of the polymerase synthesized RNA for *in vitro* selection of aptamers to human neutrophil elastase (HNE).

This invention relates to novel nucleotide triphosphate (NTP) molecules, and their incorporation into nucleic acid molecules, including nucleic acid catalysts. The NTPs of
30 the instant invention are distinct from other NTPs known in the art. The invention further relates to incorporation of these nucleotide triphosphates into oligonucleotides using an RNA polymerase; the invention further relates to novel transcription conditions for the

incorporation of modified (non-standard) and unmodified NTP's, into nucleic acid molecules. Further, the invention relates to methods for synthesis of novel NTP's

In a first aspect, the invention features NTP's having the formula triphosphate-OR, for example the following formula 3:



5

where R is any nucleoside; specifically the nucleosides 2'-O-methyl-2,6-diaminopurine riboside; 2'-deoxy-2'-amino-2,6-diaminopurine riboside; 2'-(*N*-alanyl) amino-2'-deoxy-uridine; 2'-(*N*-phenylalanyl) amino-2'-deoxy-uridine; 2'-deoxy-2'-(*N*-β-alanyl) amino ; 2'-deoxy-2'-(lysiyl) amino uridine; 2'-*C*-allyl uridine; 2'-*O*-amino-uridine; 2'-*O*-methylthiomethyl adenosine; 2'-*O*-methylthiomethyl cytidine ; 2'-*O*-methylthiomethyl guanosine; 2'-*O*-methylthiomethyl-uridine; 2'-deoxy-2'-(*N*-histidyl) amino uridine; 2'-deoxy-2'-amino-5-methyl cytidine; 2'-(*N*-β-carboxamidine-β-alanyl) amino-2'-deoxy-uridine; 2'-deoxy-2'-(*N*-β-alanyl)-guanosine; 2'-*O*-amino-adenosine; 2'-(*N*-lysiyl) amino-2'-deoxy-cytidine; 2'-Deoxy-2'-(*L*-histidine) amino Cytidine; 5-Imidazoleacetic acid 2'-deoxy uridine, 5-[3-(*N*-4-imidazoleacetyl)aminopropynyl]-2'-*O*-methyl uridine, 5-(3-aminopropynyl)-2'-*O*-methyl uridine, 5-(3-aminopropyl)-2'-*O*-methyl uridine, 5-[3-(*N*-4-imidazoleacetyl)aminopropyl]-2'-*O*-methyl uridine, 5-(3-aminopropyl)-2'-deoxy-2-fluoro uridine, 2'-Deoxy-2'-(β-alanyl-*L*-histidyl) amino uridine, 2'-deoxy-2'-β-alaninamido-uridine, 3-(2'-deoxy-2'-fluoro-β-*D*-ribofuranosyl)piperazino[2,3-*D*]pyrimidine-2-one, 5-[3-(*N*-4-imidazoleacetyl)aminopropyl]-2'-deoxy-2'-fluoro uridine, 5-[3-(*N*-4-imidazoleacetyl)aminopropynyl]-2'-deoxy-2'-fluoro uridine, 5-*E*-(2-carboxyvinyl)-2'-deoxy-2'-fluoro uridine, 5-[3-(*N*-4-aspartyl)aminopropynyl]-2'-fluoro uridine, 5-(3-aminopropyl)-2'-deoxy-2-fluoro cytidine, and 5-[3-(*N*-4-succinyl)aminopropyl]-2'-deoxy-2-fluoro cytidine.

5

In a second aspect, the invention features inorganic and organic salts of the nucleoside triphosphates of the instant invention.

In a third aspect, the invention features a process for the synthesis of pyrimidine nucleotide triphosphate (such as UTP, 2'-*O*-MTM-UTP, dUTP and the like) including the steps of monophosphorylation where the pyrimidine nucleoside is contacted with a mixture having a phosphorylating agent (such as phosphorus oxychloride, phospho-tris-
5 triazolides, phospho-tris-triimidazolides and the like), trialkyl phosphate (such as triethylphosphate or trimethylphosphate or the like) and a hindered base (such as dimethylaminopyridine, DMAP and the like) under conditions suitable for the formation of pyrimidine monophosphate; and pyrophosphorylation where the pyrimidine monophosphate is contacted with a pyrophosphorylating reagent (such as
10 tributylammonium pyrophosphate) under conditions suitable for the formation of pyrimidine triphosphates.

By "nucleotide triphosphate" or "NTP" is meant a nucleoside bound to three inorganic phosphate groups at the 5' hydroxyl group of the modified or unmodified ribose or deoxyribose sugar where the 1' position of the sugar may comprise a nucleic acid base
15 or hydrogen. The triphosphate portion may be modified to include chemical moieties which do not destroy the functionality of the group (*i.e.*, allow incorporation into an RNA molecule).

In another preferred embodiment, nucleotide triphosphates (NTPs) of the instant invention are incorporated into an oligonucleotide using an RNA polymerase enzyme.
20 RNA polymerases include but are not limited to mutated and wild type versions of bacteriophage T7, SP6, or T3 RNA polymerases. Applicant has also found that the NTPs of the present invention can be incorporated into oligonucleotides using certain DNA polymerases, such as Taq polymerase.

In yet another preferred embodiment, the invention features a process for
25 incorporating modified NTP's into an oligonucleotide including the step of incubating a mixture having a DNA template, RNA polymerase, NTP, and an enhancer of modified NTP incorporation under conditions suitable for the incorporation of the modified NTP into the oligonucleotide.

By "enhancer of modified NTP incorporation" is meant a reagent which facilitates
30 the incorporation of modified nucleotides into a nucleic acid transcript by an RNA polymerase. Such reagents include, but are not limited to, methanol, LiCl, polyethylene glycol (PEG), diethyl ether, propanol, methyl amine, ethanol, and the like.

In another preferred embodiment, the modified nucleotide triphosphates can be incorporated by transcription into a nucleic acid molecules including enzymatic nucleic acid, antisense, 2-5A antisense chimera, oligonucleotides, triplex forming oligonucleotide (TFO), aptamers and the like (Stull *et al.*, 1995 *Pharmaceutical Res.* 12, 465).

5 By "triplex forming oligonucleotides (TFO)" it is meant an oligonucleotide that can bind to a double-stranded DNA in a sequence-specific manner to form a triple-strand helix. Formation of such triple helix structure has been shown to inhibit transcription of the targeted gene (Duval-Valentin *et al.*, 1992 *Proc. Natl. Acad. Sci. USA* 89, 504).

0 In yet another preferred embodiment, the modified nucleotide triphosphates of the instant invention can be used for combinatorial chemistry or *in vitro* selection of nucleic acid molecules with novel function. Modified oligonucleotides can be enzymatically synthesized to generate libraries for screening.

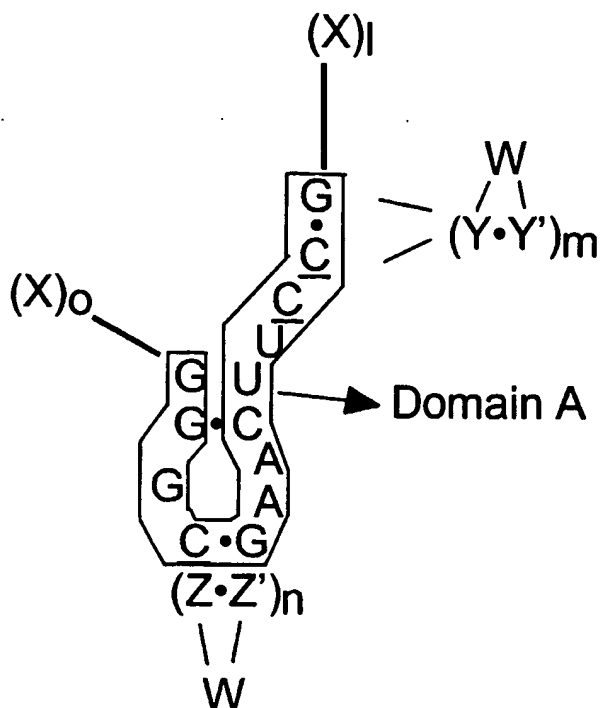
In another preferred embodiment, the invention features nucleic acid based techniques (*e.g.*, enzymatic nucleic acid molecules), antisense nucleic acids, 2-5A
5 antisense chimeras, triplex DNA, antisense nucleic acids containing RNA cleaving chemical groups) isolated using the methods described in this invention and methods for their use to diagnose, down regulate or inhibit gene expression.

In yet another preferred embodiment, the invention features enzymatic nucleic acid molecules targeted against HER2 RNA, specifically including ribozymes in the class II
10 (zinzyme) motif.

Targets, for example HER2 RNA, for useful ribozymes and antisense nucleic acids can be determined, for example, as described in Draper *et al.*, WO 93/23569; Sullivan *et al.*, WO 93/23057; Thompson *et al.*, WO 94/02595; Draper *et al.*, WO 95/04818; McSwiggen *et al.*, US Patent Nos. 5,525,468 and 5,646,042, all are hereby incorporated by
5 reference herein in their totalities. Other examples include the following PCT applications, which concern inactivation of expression of disease-related genes: WO 95/23225, and WO 95/13380; all of which are incorporated by reference herein.

In yet another preferred embodiment, the invention features a process for incorporating a plurality of compounds of formula 3.

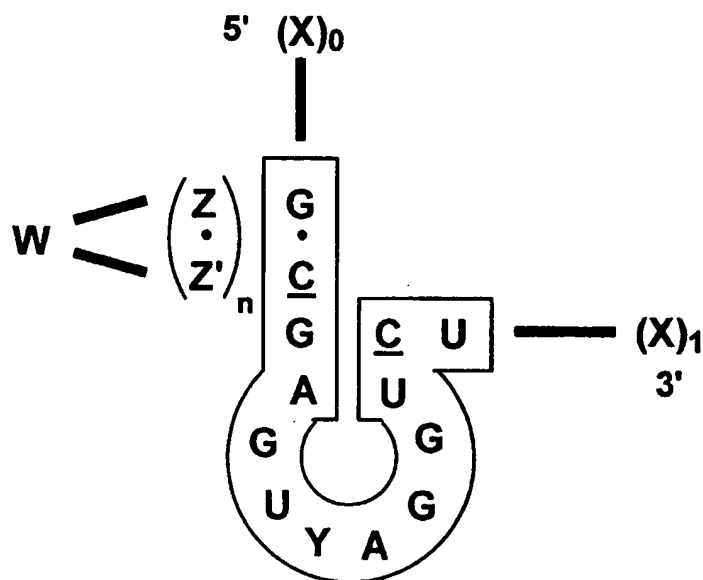
0 In yet another embodiment, the invention features a nucleic acid molecule with catalytic activity having formula 4:



- In the formula shown above X, Y, and Z represent independently a nucleotide or a non-nucleotide linker, which may be same or different; • indicates hydrogen bond formation between two adjacent nucleotides which may or may not be present; Y' is a nucleotide complementary to Y; Z' is a nucleotide complementary to Z; l is an integer greater than or equal to 3 and preferably less than 20, more specifically 4, 5, 6, 7, 8, 9, 10, 11, 12, or 15; m is an integer greater than 1 and preferably less than 10, more specifically 2, 3, 4, 5, 6, or 7; n is an integer greater than 1 and preferably less than 10, more specifically 3, 4, 5, 6, or 7; o is an integer greater than or equal to 3 and preferably less than 20, more specifically 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, or 15; l and o may be the same length ($l = o$) or different lengths ($l \neq o$); each $X(l)$ and $X(o)$ are oligonucleotides which are of sufficient length to stably interact independently with a target nucleic acid sequence (the target can be an RNA, DNA or RNA/DNA mixed polymers); W is a linker of ≥ 2 nucleotides in length or may be a non-nucleotide linker; A, U, C, and G represent the nucleotides; G is a nucleotide, preferably 2'-O-methyl or ribo; A is a nucleotide, preferably 2'-O-methyl or ribo; U is a nucleotide, preferably 2'-amino (e.g., 2'-NH₂ or 2'-O-NH₂), 2'-O-methyl or ribo; C represents a nucleotide, preferably 2'-amino (e.g., 2'-NH₂ or 2'-O-

NH₂), and _____ represents a chemical linkage (e.g. a phosphate ester linkage, amide linkage, phosphorothioate, phosphorodithioate or others known in the art).

In yet another embodiment, the invention features a nucleic acid molecule with catalytic activity having formula 5:



- 5 In the formula shown above X, Y, and Z represent independently a nucleotide or a non-nucleotide linker, which may be same or different; • indicates hydrogen bond formation between two adjacent nucleotides which may or may not be present; Z' is a nucleotide complementary to Z; l is an integer greater than or equal to 3 and preferably less than 20, more specifically 4, 5, 6, 7, 8, 9, 10, 11, 12, or 15; n is an integer greater than
- 10 1 and preferably less than 10, more specifically 3, 4, 5, 6, or 7; o is an integer greater than or equal to 3 and preferably less than 20, more specifically 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, or 15; l and o may be the same length (l = o) or different lengths (l ≠ o); each X_(l) and X_(o) are oligonucleotides which are of sufficient length to stably interact independently with a target nucleic acid sequence (the target can be an RNA, DNA or RNA/DNA mixed
- 15 polymers); X_(o) preferably has a G at the 3'-end, X_(l) preferably has a G at the 5'-end; W is a linker of ≥ 2 nucleotides in length or may be a non-nucleotide linker; Y is a linker of ≥ 1 nucleotides in length, preferably G, 5'-CA-3', or 5'-CAA-3', or may be a non-nucleotide linker; A, U, C, and G represent nucleotides; G is a nucleotide, preferably 2'-O-methyl, 2'-deoxy-2'-fluoro, or 2'-OH; A is a nucleotide, preferably 2'-O-methyl, 2'-deoxy-2'-

fluoro, or 2'-OH; U is a nucleotide, preferably 2'-O-methyl, 2'-deoxy-2'-fluoro, or 2'-OH; C represents a nucleotide, preferably 2'-amino (e.g., 2'-NH₂ or 2'-O- NH₂, and _____ represents a chemical linkage (e.g. a phosphate ester linkage, amide linkage, phosphorothioate, phosphorodithioate or others known in the art).

5 The enzymatic nucleic acid molecules of Formula 4 and Formula 5 may independently comprise a cap structure which may independently be present or absent.

 In yet another preferred embodiment, the 3'-cap is selected from a group comprising, 4',5'-methylene nucleotide; 1-(beta-D-erythrofuransyl) nucleotide; 4'-thio nucleotide; carbocyclic nucleotide; 5'-amino-alkyl phosphate; 1,3-diamino-2-propyl phosphate; 3-aminopropyl phosphate; 6-aminohexyl phosphate; 1,2-aminododecyl phosphate; hydroxypropyl phosphate; 1,5-anhydrohexitol nucleotide; L-nucleotide; alpha-nucleotide; modified base nucleotide; phosphorodithioate; *threo*-pentofuransyl nucleotide; acyclic 3',4'-seco nucleotide; 3,4-dihydroxybutyl nucleotide; 3,5-dihydroxypentyl nucleotide; 5'-5'-inverted nucleotide moiety; 5'-5'-inverted abasic moiety; 5'-phosphoramidate; 5'-phosphorothioate; 1,4-butanediol phosphate 5'-amino; bridging and/or non-bridging 5'-phosphoramidate, phosphorothioate and/or phosphorodithioate; bridging or non bridging methylphosphonate and 5'-mercapto moieties (for more details, see Beaucage and Iyer, 1993, *Tetrahedron* 49, 1925; incorporated by reference herein).

15 In another aspect, the invention provides mammalian cells containing one or more nucleic acid molecules and/or expression vectors of this invention. The one or more nucleic acid molecules may independently be targeted to the same or different sites.

Nucleotide Synthesis

 Addition of dimethylaminopyridine (DMAP) to the phosphorylation protocols known in the art can greatly increase the yield of nucleotide monophosphates while decreasing the reaction time. Synthesis of the nucleosides of the invention have been described in several publications and Applicants previous applications (Beigelman *et al.*, International PCT publication No. WO 96/18736; Dudzcy *et al.*, Int. PCT Pub. No. WO 95/11910; Usman *et al.*, Int. PCT Pub. No. WO 95/13378; Matulic-Adamic *et al.*, 1997, *Tetrahedron Lett.* 38, 203; Matulic-Adamic *et al.*, 1997, *Tetrahedron Lett.* 38, 1669; all of which are incorporated herein by reference). These nucleosides are dissolved in triethyl phosphate and chilled in an ice bath. Phosphorus oxychloride (POCl₃) is then added

followed by the introduction of DMAP. The reaction is then warmed to room temperature and allowed to proceed for 5 hours. This reaction allows the formation of nucleotide monophosphates which can then be used in the formation of nucleotide triphosphates. Tributylamine is added followed by the addition of anhydrous acetonitrile and
5 tributylammonium pyrophosphate. The reaction is then quenched with TEAB and stirred overnight at room temperature (about 20°C). The triphosphate is purified using Sephadex® column purification or equivalent and/or HPLC and the chemical structure is confirmed using NMR analysis. Those skilled in the art will recognize that the reagents, temperatures of the reaction, and purification methods can easily be alternated with
10 substitutes and equivalents and still obtain the desired product.

Nucleotide Triphosphates

The invention provides nucleotide triphosphates which can be used for a number of different functions. The nucleotide triphosphates formed from nucleosides found in Table
15 45 are unique and distinct from other nucleotide triphosphates known in the art. Incorporation of modified nucleotides into DNA or RNA oligonucleotides can alter the properties of the molecule. For example, modified nucleotides can hinder binding of nucleases, thus increasing the chemical half-life of the molecule. This is especially important if the molecule is to be used for cell culture or *in vivo*. It is known in the art that
20 the introduction of modified nucleotides into these molecules can greatly increase the stability and thereby the effectiveness of the molecules (Burgin *et al.*, 1996, *Biochemistry* 35, 14090-14097; Usman *et al.*, 1996, *Curr. Opin. Struct. Biol.* 6, 527-533).

Modified nucleotides are incorporated using either wild type or mutant polymerases. For example, mutant T7 polymerase is used in the presence of modified nucleotide
25 triphosphate(s), DNA template and suitable buffers. Those skilled in the art will recognize that other polymerases and their respective mutant versions can also be utilized for the incorporation of NTP's of the invention. Nucleic acid transcripts were detected by incorporating radiolabelled nucleotides (α -³²P NTP). The radiolabeled NTP contained the same base as the modified triphosphate being tested. The effects of methanol, PEG and
30 LiCl were tested by adding these compounds independently or in combination. Detection and quantitation of the nucleic acid transcripts was performed using a Molecular Dynamics

PhosphorImager. Efficiency of transcription was assessed by comparing modified nucleotide triphosphate incorporation with all-ribonucleotide incorporation control. Wild-type polymerase was used to incorporate NTP's using the manufacturer's buffers and instructions (Boehringer Mannheim).

5

Transcription Conditions

Incorporation rates of modified nucleotide triphosphates into oligonucleotides can be increased by adding to traditional buffer conditions, several different enhancers of modified NTP incorporation. Applicant has utilized methanol and LiCl in an attempt to increase incorporation rates of dNTP using RNA polymerase. These enhancers of modified NTP incorporation can be used in different combinations and ratios to optimize transcription. Optimal reaction conditions differ between nucleotide triphosphates and can readily be determined by standard experimentation. Overall, however, Applicant has found that inclusion of enhancers of modified NTP incorporation such as methanol or inorganic compound such as lithium chloride increase the mean transcription rates.

15

Applicant synthesized pyrimidine nucleotide triphosphates using DMAP in the reaction. For purines, applicant utilized standard protocols previously described in the art (Yoshikawa *et al supra*; Ludwig, *supra*). Described below is one example of a pyrimidine nucleotide triphosphate and one purine nucleotide triphosphate synthesis.

20

Synthesis of purine nucleotide triphosphates: 2'-O-methyl-guanosine-5'-triphosphate

2'-O-methyl guanosine nucleoside (0.25 grams, 0.84 mmol) was dissolved in triethyl phosphate (5.0) ml by heating to 100°C for 5 minutes. The resulting clear, colorless solution was cooled to 0°C using an ice bath under an argon atmosphere. Phosphorous oxychloride (1.8 eq., 0.141 ml) was then added to the reaction mixture with vigorous stirring. The reaction was monitored by HPLC, using a sodium perchlorate gradient. After 5 hours at 0°C, tributylamine (0.65 ml) was added followed by the addition of anhydrous acetonitrile (10.0 ml), and after 5 minutes (reequilibration to 0°C) tributylammonium pyrophosphate (4.0 eq., 1.53 g) was added. The reaction mixture was quenched with 20 ml of 2M TEAB after 15 minutes at 0°C (HPLC analysis with above conditions showed consumption of monophosphate at 10 minutes) then stirred overnight at room temperature, the mixture was evaporated *in vacuo* with methanol co-evaporation

25

30

(4x) then diluted in 50 ml 0.05M TEAB. DEAE sephadex purification was used with a gradient of 0.05 to 0.6 M TEAB to obtain pure triphosphate (0.52 g, 66.0% yield) (elutes around 0.3M TEAB); the purity was confirmed by HPLC and NMR analysis.

5 Synthesis of Pyrimidine nucleotide triphosphates: 2'-O-methylthiomethyl-uridine-5'-triphosphate

2'-O-methylthiomethyl uridine nucleoside (0.27 grams, 1.0 mmol) was dissolved in triethyl phosphate (5.0 ml). The resulting clear, colorless solution was cooled to 0°C with an ice bath under an argon atmosphere. Phosphorus oxychloride (2.0 eq., 0.190 ml) was
10 then added to the reaction mixture with vigorous stirring. Dimethylaminopyridine (DMAP, 0.2eq., 25 mg) was added, the solution warmed to room temperature and the reaction was monitored by HPLC, using a sodium perchlorate gradient. After 5 hours at 20°C, tributylamine (1.0 ml) was added followed by anhydrous acetonitrile (10.0 ml), and after 5 minutes tributylammonium pyrophosphate (4.0 eq., 1.8 g) was added. The reaction
15 mixture was quenched with 20 ml of 2M TEAB after 15 minutes at 20°C (HPLC analysis with above conditions showed consumption of monophosphate at 10 minutes) then stirred overnight at room temperature. The mixture was evaporated *in vacuo* with methanol co-evaporation (4x) then diluted in 50 ml 0.05M TEAB. DEAE fast flow Sepharose purification with a gradient of 0.05 to 1.0 M TEAB was used to obtain pure triphosphate
20 (0.40 g, 44% yield) (elutes around 0.3M TEAB) as determined by HPLC and NMR analysis.

Utilization of DMAP in Uridine 5'-Triphosphate Synthesis

The reactions were performed on 20 mg aliquots of nucleoside dissolved in 1 ml of
25 triethyl phosphate and 19 ul of phosphorus oxychloride. The reactions were monitored at 40 minute intervals automatically by HPLC to generate yield-of-product curves at times up to 18 hours. A reverse phase column and ammonium acetate/ sodium acetate buffer system (50mM & 100mM respectively at pH 4.2) was used to separate the 5', 3', 2' monophosphates (the monophosphates elute in that order) from the 5'-triphosphate and the
30 starting nucleoside. The data is shown in Table 46. These conditions doubled the product yield and resulted in a 10-fold improvement in the reaction time to maximum yield (1200 minutes down to 120 minutes for a 90% yield). Selectivity for 5'-monophosphorylation

was observed for all reactions. Subsequent triphosphorylation occurred in nearly quantitative yield.

Materials Used in Bacteriophage T7 RNA Polymerase Reactions

5 **Buffer 1:** Reagents are mixed together to form a 10X stock solution of buffer 1 (400 mM Tris-Cl [pH 8.1], 200 mM MgCl₂, 100 mM DTT, 50 mM spermidine, and 0.1% triton® X-100). Prior to initiation of the polymerase reaction methanol, LiCl is added and the buffer is diluted such that the final reaction conditions for condition 1 consisted of :
40mM tris (pH 8.1), 20mM MgCl₂, 10 mM DTT, 5 mM spermidine, 0.01% triton® X-
10 100, 10% methanol, and 1 mM LiCl.

BUFFER 2: Reagents are mixed together to form a 10X stock solution of buffer 2 (400 mM Tris-Cl [pH 8.1], 200 mM MgCl₂, 100 mM DTT, 50 mM spermidine, and 0.1% triton® X-100). Prior to initiation of the polymerase reaction PEG, LiCl is added and the buffer is diluted such that the final reaction conditions for buffer 2 consisted of : 40mM
15 tris (pH 8.1), 20mM MgCl₂, 10 mM DTT, 5 mM spermidine, 0.01% triton® X-100, 4% PEG, and 1 mM LiCl.

BUFFER 3: Reagents are mixed together to form a 10X stock solution of buffer 3 (400 mM Tris-Cl [pH 8.0], 120 mM MgCl₂, 50 mM DTT, 10 mM spermidine and 0.02% triton® X-100). Prior to initiation of the polymerase reaction PEG is added and the buffer
20 is diluted such that the final reaction conditions for buffer 3 consisted of : 40mM tris (pH 8.0), 12 mM MgCl₂, 5 mM DTT, 1 mM spermidine, 0.002% triton® X-100, and 4% PEG.

BUFFER 4: Reagents are mixed together to form a 10X stock solution of buffer 4 (400 mM Tris-Cl [pH 8.0], 120 mM MgCl₂, 50 mM DTT, 10 mM spermidine and 0.02% triton® X-100). Prior to initiation of the polymerase reaction PEG, methanol is added and
25 the buffer is diluted such that the final reaction conditions for buffer 4 consisted of : 40mM tris (pH 8.0), 12 mM MgCl₂, 5 mM DTT, 1 mM spermidine, 0.002% triton® X-100, 10% methanol, and 4% PEG.

BUFFER 5: Reagents are mixed together to form a 10X stock solution of buffer 5 (400 mM Tris-Cl [pH 8.0], 120 mM MgCl₂, 50 mM DTT, 10 mM spermidine and 0.02% triton® X-100). Prior to initiation of the polymerase reaction PEG, LiCl is added and the
30 buffer is diluted such that the final reaction conditions for buffer 5 consisted of : 40mM

tris (pH 8.0), 12 mM MgCl₂, 5 mM DTT, 1 mM spermidine, 0.002% triton® X-100, 1 mM LiCl and 4% PEG.

BUFFER 6: Reagents are mixed together to form a 10X stock solution of buffer 6 (400 mM Tris-Cl [pH 8.0], 120 mM MgCl₂, 50 mM DTT, 10 mM spermidine and 0.02% triton® X-100). Prior to initiation of the polymerase reaction PEG, methanol is added and the buffer is diluted such that the final reaction conditions for buffer 6 consisted of :
40mM tris (pH 8.0), 12 mM MgCl₂, 5 mM DTT, 1 mM spermidine, 0.002% triton® X-100, 10% methanol, and 4% PEG.

BUFFER 7: Reagents are mixed together to form a 10X stock solution of buffer 6 (400 mM Tris-Cl [pH 8.0], 120 mM MgCl₂, 50 mM DTT, 10 mM spermidine and 0.02% triton® X-100). Prior to initiation of the polymerase reaction PEG, methanol and LiCl is added and the buffer is diluted such that the final reaction conditions for buffer 6 consisted of : 40mM tris (pH 8.0), 12 mM MgCl₂, 5 mM DTT, 1 mM spermidine, 0.002% triton® X-100, 10% methanol, 4% PEG, and 1 mM LiCl.

15

Screening of Modified nucleotide triphosphates with Mutant T7 RNA Polymerase

Modified nucleotide triphosphates were tested in buffers 1 through 6 at two different temperatures (25 and 37°C). Buffers 1-6 tested at 25°C were designated conditions 1-6 and buffers 1-6 tested at 37°C were designated conditions 7-12 (**Table 47**). In each condition, Y639F mutant T7 polymerase (Sousa and Padilla, *supra*) (0.3-2 mg/20 ml reaction), NTP's (2 mM each), DNA template (10 pmol), inorganic pyrophosphatase (5U/ml) and α -³²P NTP (0.8 mCi/pmol template) were combined and heated at the designated temperatures for 1-2 hours. The radiolabeled NTP used was different from the modified triphosphate being testing. The samples were resolved by polyacrylamide gel electrophoresis. Using a PhosphorImager (Molecular Dynamics, Sunnyvale, CA), the amount of full-length transcript was quantified and compared with an all-RNA control reaction. The data is presented in **Table 48**; results in each reaction are expressed as a percent compared to the all-ribonucleotide triphosphate (rNTP) control. The control was run with the mutant T7 polymerase using commercially available polymerase buffer (Boehringer Mannheim, Indianapolis, IN).

30

Incorporation of Modified NTP's using Wild-type T7 RNA polymerase

Bacteriophage T7 RNA polymerase was purchased from Boehringer Mannheim at 0.4 U/ μ L concentration. Applicant used the commercial buffer supplied with the enzyme and 0.2 μ Ci alpha- 32 P NTP in a 50 μ L reaction with nucleotides triphosphates at 2 mM each. The template was a double-stranded PCR fragment, which was used in previous screens. Reactions were carried out at 37°C for 1 hour. Ten μ L of the sample was run on a 7.5% analytical PAGE and bands were quantitated using a PhosphorImager. Results are calculated as a comparison to an "all ribo" control (non-modified nucleotide triphosphates) and the results are in **Table 49**.

10 Incorporation of Multiple Modified nucleotide triphosphates Into Oligonucleotides

Combinations of modified nucleotide triphosphates were tested with the transcription protocol described above, to determine the rates of incorporation of two or more of these triphosphates. Incorporation of 2'-Deoxy-2'-(L-histidine) amino uridine (2'-his-NH₂-UTP) was tested with unmodified cytidine nucleotide triphosphates, rATP and rGTP in reaction condition number 9. The data is presented as a percentage of incorporation of modified NTP's compared to the all rNTP control and is shown in **Table 50a**.

Two modified cytidines (2'-NH₂-CTP or 2'dCTP) were incorporated along with 2'-his-NH₂-UTP with identical efficiencies. 2'-his-NH₂-UTP and 2'-NH₂-CTP were then tested with various unmodified and modified adenosine triphosphates in the same buffer (**Table 50b**). The best modified adenosine triphosphate for incorporation with both 2'-his-NH₂-UTP and 2'-NH₂-CTP was 2'-NH₂-DAPTP.

25 Optimization of Reaction conditions for Incorporation of Modified Nucleotide Triphosphate

The combination of 2'-his-NH₂-UTP, 2'-NH₂-CTP, 2'-NH₂-DAP, and rGTP was tested in several reaction conditions (**Table 51**) using the incorporation protocol described above. The results demonstrate that of the buffer conditions tested, incorporation of these modified nucleotide triphosphates occur in the presence of both methanol and LiCl.

Selection of Novel Enzymatic nucleic acid molecule Motifs using 2'-deoxy-2' amino
Modified GTP and CTP

For selection of new enzymatic nucleic acid molecule motifs, pools of enzymatic nucleic acid molecules were designed to have two substrate binding arms (5 and 16 nucleotides long) and a random region in the middle. The substrate has a biotin on the 5' end, 5 nucleotides complementary to the short binding arm of the pool, an unpaired G (the desired cleavage site), and 16 nucleotides complementary to the long binding arm of the pool. The substrate was bound to column resin through an avidin-biotin complex. The general process for selection is shown in **Figure 11**. The protocols described below represent one possible method that may be utilized for selection of enzymatic nucleic acid molecules and are given as a non-limiting example of enzymatic nucleic acid molecule selection with combinatorial libraries.

Construction of Libraries:

The oligonucleotides listed below were synthesized by Operon Technologies (Alameda, CA). Templates were gel purified and then run through a Sep-Pak™ cartridge (Waters, Millford, MA) using the manufacturers protocol. Primers (MST3, MST7c, MST3del) were used without purification.

Primers:

MST3 (30 mer): 5'-CAC TTA GCA TTA ACC CTC ACT AAA GGC CGT-3'
MST7c (33 mer): 5'-TAA TAC GAC TCA CTA TAG GAA AGG TGT GCA ACC-3'
MST3del (18 mer): 5'-ACC CTC ACT AAA GGC CGT-3'

Templates:

MSN60c (93 mer): 5'-ACC CTC ACT AAA GGC CGT (N)₆₀ GGT TGC ACA CCT
TTG-3'
MSN40c (73 mer): 5'-ACC CTC ACT AAA GGC CGT (N)₄₀ GGT TGC ACA CCT
TTG-3'
MSN20c (53 mer): 5'-ACC CTC ACT AAA GGC CGT (N)₂₀ GGT TGC ACA CCT
TTG-3'

N60 library was constructed using MSN60c as a template and MST3/MST7c as primers. N40 and N20 libraries were constructed using MSN40c (or MSN20c) as template and MST3del/MST7c as primers.

Single-stranded templates were converted into double-stranded DNA by the following protocol: 5 nmol template, 10 nmol each primer, in 10 ml reaction volume using standard PCR buffer, dNTP's, and taq DNA polymerase (all reagents from Boehringer Mannheim). Synthesis cycle conditions were 94°C, 4 minutes; (94°C, 1
5 minute; 42°C, 1 minute; 72°C, 2 minutes) x 4; 72°C, 10 minutes. Products were checked on agarose gel to confirm the length of each fragment (N60=123 bp, N40=91 bp, N20=71 bp) and then were phenol/chloroform extracted and ethanol precipitated. The concentration of the double-stranded product was 25 µM.

Transcription of the initial pools was performed in a 1 ml volume comprising: 500
10 pmol double-stranded template (3×10^{14} molecules), 40 mM tris-HCl (pH 8.0), 12 mM MgCl₂, 1 mM spermidine, 5 mM DTT, 0.002% triton X-100, 1 mM LiCl, 4% PEG 8000, 10% methanol, 2 mM ATP (Pharmacia), 2 mM GTP (Pharmacia), 2 mM 2'-deoxy-2'-amino-CTP (USB), 2 mM 2'-deoxy-2'-amino-UTP (USB), 5 U/ml inorganic pyrophosphatase (Sigma), 5 U/µl T7 RNA polymerase (USB; Y639F mutant was used in
15 some cases at 0.1 mg/ml (Sousa and Padilla, *supra*)), 37°C, 2 hours. Transcribed libraries were purified by denaturing PAGE (N60=106 ntds, N40=74, N20=54) and the resulting product was desalted using Sep-Pak™ columns and then ethanol precipitated.

Initial column-Selection:

20 The following biotinylated substrate was synthesized using standard protocols (Usman *et al.*, 1987 *J. Am. Chem. Soc.*, 109, 7845; Scaringe *et al.*, 1990 *Nucleic Acids Res.*, 18, 5433; and Wincott *et al.*, 1995 *Nucleic Acids Res.* 23, 2677-2684):

5'-biotin-C18 spacer-GCC GUG GGU UGC ACA CCU UUC C-C18 spacer-thiol-modifier C6 S-S-inverted abasic-3'

25 Substrate was purified by denaturing PAGE and ethanol precipitated. 10 nmol of substrate was linked to a NeutrAvidin™ column using the following protocol: 400 µl UltraLink Immobilized NeutrAvidin™ slurry (200 µl beads, Pierce, Rockford, IL) were loaded into a polystyrene column (Pierce). The column was washed twice with 1 ml of binding buffer (20 mM NaPO₄ (pH 7.5), 150 mM NaCl) and then capped off (i.e., a cap
30 was put on the bottom of the column to stop the flow). 200 µl of the substrate suspended in binding buffer was applied and allowed to incubate at room temperature for 30 minutes

with occasional vortexing to ensure even linking and distribution of the solution to the resin. After the incubation, the cap was removed and the column was washed with 1 ml binding buffer followed by 1 ml column buffer (50 mM tris-HCL (pH 8.5), 100 mM NaCl, 50 mM KCl). The column was then ready for use and capped off. 1 nmol of the initial pool RNA was loaded on the column in a volume of 200 µl column buffer. It was allowed to bind the substrate by incubating for 30 minutes at room temperature with occasional vortexing. After the incubation, the cap was removed and the column was washed twice with 1 ml column buffer and capped off. 200 µl of elution buffer (50 mM tris-HCL (pH 8.5), 100 mM NaCl, 50 mM KCl, 25 mM MgCl₂) was applied to the column followed by 30 minute incubation at room temperature with occasional vortexing. The cap was removed and four 200 µl fractions were collected using elution buffer.

Second column (counter selection):

A diagram for events in the second column is generally shown in Figure 12 and substrate oligonucleotide used is shown below:

5'-GGU UGC ACA CCU UUC C-C18 spacer-biotin-inverted abasic-3'

This column substrate was linked to UltraLink NeutrAvidin™ resin as previously described (40 pmol) which was washed twice with elution buffer. The eluent from the first column purification was then run on the second column. The use of this column allowed for binding of RNA that non-specifically diluted from the first column, while RNA that performed a catalytic event and had product bound to it, flowed through the second column. The fractions were ethanol precipitated using glycogen as carrier and rehydrated in sterile water for amplification.

Amplification:

RNA and primer MST3 (10-100 pmol) were denatured at 90°C for 3 minutes in water and then snap-cooled on ice for one minute. The following reagents were added to the tube (final concentrations given): 1X PCR buffer (Boehringer Mannheim), 1 mM dNTP's (for PCR, Boehringer Mannheim), 2 U/µl RNase-Inhibitor (Boehringer Mannheim), 10 U/µl Superscript™ II Reverse Transcriptase (BRL). The reaction was incubated for 1 hour at 42°C, then at 95°C for 5 minutes in order to destroy the

Superscript™. The following reagents were then added to the tube to increase the volume five-fold for the PCR step (final concentrations/amounts given): MST7c primer (10-100 pmol, same amount as in RT step), 1X PCR buffer, taq DNA polymerase (0.025-0.05 U/μl, Boehringer Mannheim). The reaction was cycled as follows: 94°C, 4minutes;
5 (94°C, 30s; 42-54°C, 30s; 72°C, 1minute) x 4-30 cycles; 72°C, 5minutes; 30°C, 30 minutes. Cycle number and annealing temperature were decided on a round by round basis. In cases where heteroduplex was observed, the reaction was diluted five-fold with fresh reagents and allowed to progress through 2 more amplification cycles. Resulting products were analyzed for size on an agarose gel (N60=123 bp, N40=103 bp, N20=83 bp)
10 and then ethanol precipitated.

Transcriptions:

Transcription of amplified products was done using the conditions described above with the following modifications: 10-20% of the amplification reaction was used as
15 template, reaction volume was 100-500 μl, and the products sizes varied slightly (N60=106 ntds, N40=86, N20=66). A small amount of ³²P-GTP was added to the reactions for quantitation purposes.

Subsequent rounds:

20 Subsequent rounds of selection used 20 pmols of input RNA and 40 pmol of the 22 nucleotide substrate on the column.

Activity of pools:

Pools were assayed for activity under single turnover conditions every three to four
25 rounds. Activity assay conditions were as follows: 50 mM tris-HCl (pH 8.5), 25 mM MgCl₂, 100 mM NaCl, 50 mM KCl, trace ³²P-labeled substrate, 10 nM RNA pool. 2X pool in buffer and, separately, 2X substrate in buffer were incubated at 90°C for 3 minutes, then at 37°C for 3 minutes. Equal volume 2X substrate was then added the 2X pool tube (t=0). Initial assay time points were taken at 4 and 24 hours: 5 μl was removed and
30 quenched in 8 μl cold Stop buffer (96% formamide, 20 mM EDTA, 0.05% bromphenyl blue/xylene cyanol). Samples were heated 90°C, 3 minutes, and loaded on a 20%

sequencing gel. Quantitation was performed using a Molecular Dynamics Phosphorimager and ImageQuaNT™ software. The data is shown in Table 52.

Samples from the pools of oligonucleotide were cloned into vectors and sequenced using standard protocols (Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*,
5 Cold Spring Harbor Laboratory Press). The enzymatic nucleic acid molecules were transcribed from a representative number of these clones using methods described in this application. Individuals from each pool were tested for RNA cleavage from N60 and N40 by incubating the enzymatic nucleic acid molecules from the clones with 5/16 substrate in 2mM MgCl₂, pH 7.5, 10mM KCl at 37°C. The data in Table 54 shows that the enzymatic
10 nucleic acid molecules isolated from the pool are individually active.

Kinetic Activity:

Kinetic activity of the enzymatic nucleic acid molecule shown in Table 54, was determined by incubating enzymatic nucleic acid molecule (10 nM) with substrate in a
15 cleavage buffer (pH 8.5, 25 mM MgCl₂, 100 mM NaCl, 50 mM KCl) at 37°C.

Magnesium Dependence:

Magnesium dependence of round 15 of N20 was tested by varying MgCl₂ while other conditions were held constant (50 mM tris [pH 8.0], 100 mM NaCl, 50 mM KCl,
20 single turnover, 10 nM pool). The data is shown in Table 55, which demonstrates increased activity with increased magnesium concentrations.

Selection of Novel Enzymatic nucleic acid molecule Motifs using 2'-Deoxy-2'-(N-histidyl) amino UTP, 2'-Fluoro-ATP, and 2'-deoxy-2'-amino CTP and GTP

25 The method used for selection of novel enzymatic nucleic acid molecule motifs using 2'-deoxy-2' amino modified GTP and CTP was repeated using 2'-Deoxy-2'-(N-histidyl) amino UTP, 2'-Fluoro-ATP, and 2'-deoxy-2'-amino CTP and GTP. However, rather than causing cleavage on the initial column with MgCl₂, the initial random modified-RNA pool was loaded onto substrate-resin in the following buffer; 5 mM
30 NaOAc pH 5.2, 1 M NaCl at 4° C. After ample washing, the resin was moved to 22 ° C and the buffer switch 20 mM HEPES pH 7.4, 140 mM KCl, 10 mM NaCl, 1 mM CaCl₂, 1 mM MgCl₂. In one selection of N60 oligonucleotides, no divalent cations (MgCl₂,

CaCl₂) was used. The resin was incubated for 10 minutes to allow reaction and the eluant collected.

The enzymatic nucleic acid molecule pools were capable of cleaving 1-3% of the present substrate even in the absence of divalent cations, the background (in the absence of modified pools) was 0.2 - 0.4 %.

Synthesis of 5-substituted 2'-modified nucleosides

When designing monomeric nucleoside triphosphates for selection of therapeutic catalytic RNAs, one has to take into account nuclease stability of such molecules in biological sera. A common approach to increase RNA stability is to replace the sugar 2'-OH group with other groups like 2'-fluoro, 2'-O-methyl or 2'-amino. Fortunately such 2'-modified pyrimidine 5'triphosphates are shown to be substrates for RNA polymerases. (Aurup, H.; Williams, D.M.; Eckstein, F. *Biochemistry* 1992, 31, 9637; and Padilla, R.; Sousa, R. *Nucleic Acids Res.* 1999, 27, 1561.) On the other hand it was shown that variety of substituents at pyrimidine 5-position is well tolerated by T7 RNA polymerase (Tarasow, T.M.; Eaton, B.E. *Biopolymers* 1998, 48, 29), most likely because the natural hydrogen-bonding pattern of these nucleotides is preserved. We have chosen 2'-fluoro and 2'-O-methyl pyrimidine nucleosides as starting materials for attachment of different functionalities to the 5-position of the base. Both rigid (alkynyl) and flexible (alkyl) spacers are used. The choice of imidazole, amino and carboxylate pendant groups is based on their ability to act as general acids, general bases, nucleophiles and metal ligands, all of which can improve the catalytic effectiveness of selected nucleic acids. **Figures 21 – 24** relate to the synthesis of these compounds.

2'-O-methyluridine was 3',5'-bis-acetylated using acetic anhydride in pyridine and then converted to its 5-iodo derivative **1a** using I₂/ceric ammonium nitrate reagent (Asakura, J.; Robins, M.J. *J. Org. Chem.* 1990, 55, 4928) (Scheme 1). Both reactions proceeded in a quantitative yield and no chromatographic purifications were needed. Coupling between **1** and *N*-trifluoroacetyl propargylamine using copper(I) iodide and tetrakis(triphenylphosphine)palladium(0) catalyst as described by Hobbs (Hobbs, F.W., Jr. *J. Org. Chem.* 1989, 54, 3420) yielded **2a** in 89% yield. Selective *O*-deacetylation with aqueous NaOH afforded **3a** which was phosphorylated with POCl₃/triethylphosphate

(TEP) in the presence of 1,8-bis(dimethylamino)naphthalene (Proton-Sponge) (Method A) (Kovácz, T; Ötvös, L. Tetrahedron Lett. 1988, 29, 4525). The intermediate nucleoside phosphorodichloridate was condensed *in situ* with tri-*n*-butylammonium pyrophosphate. At the end, the *N*-TFA group was removed with concentrated ammonia. 5'-Triphosphate
5 was purified on Sephadex® DEAE A-25 ion exchange column using a linear gradient of 0.1-0.8M triethylammonium bicarbonate (TEAB) for elution. Traces of contaminating inorganic pyrophosphate are removed using C-18 RP HPLC to afford analytically pure material. Conversion into Na-salt was achieved by passing the aqueous solution of triphosphate through Dowex 50WX8 ion exchange resin in Na⁺ form to afford 4a in 45%
10 yield. When Proton-Sponge was omitted in the first phosphorylation step, yields were reduced to 10-20%. Catalytic hydrogenation of 3a yielded 5-aminopropyl derivative 5a which was phosphorylated under conditions identical to those described for propynyl derivative 3a to afford triphosphate 6a in 50% yield.

For the preparation of imidazole derivatized triphosphates 9a and 11a, we
15 developed an efficient synthesis of *N*-diphenylcarbamoyl 4-imidazoleacetic acid (ImAA^{DPC}): Transient protection of carboxyl group as TMS-ester using TMS-Cl/pyridine followed by DPC-Cl allowed for a clean and quantitative conversion of 4-imidazoleacetic acid (ImAA) to its *N*-DPC protected derivative.

Complete deacylation of 2a afforded 5-(3-aminopropynyl) derivative 8a which
20 was condensed with 4-imidazoleacetic acid in the presence of 1-(3-dimethylaminopropyl)-3-ethylcarbodiimide (EDC) to afford 9a in 68% yield. Catalytic hydrogenation of 8a yielded 5-(3-aminopropyl) derivative 10a which was condensed with ImAA^{DPC} to yield conjugate 11a in 32% yield. Yields in these couplings were greatly improved when 5'-OH was protected with DMT group (not shown) thus
25 efficiently preventing undesired 5'-*O*-esterification. Both 9a and 11a failed to yield triphosphate products in reaction with POCl₃/TEP/Proton-Sponge.

On the contrary, phosphorylation of 3'-*O*-acetylated derivatives 12a and 13a using 2-chloro-4*H*-1,3,2-benzodioxaphosphorin-4-one followed by pyrophosphate addition and oxidation (Method B, Scheme 2; Ludwig, J., Eckstein, F., *J. Org. Chem.* 1989, 54,
30 631) afforded the desired triphosphates 14a and 15a in 57% yield, respectively.

2'-Deoxy-2'-fluoro nucleoside 5'-triphosphates containing amino- (**4b**, **6b**) and imidazole- (**14b**, **15b**) linked groups were synthesized in a manner analogous to that described for the preparation of 2'-*O*-methyl nucleoside 5'-triphosphates (Schemes 1 and 2). Again, only Ludwig-Eckstein's phosphorylation worked for the preparation of 4-imidazoleacetyl derivatized triphosphates.

It is worth noting that when "one-pot-two-steps" phosphorylation reaction (Kovácz, T; Ötvös, L. *Tetrahedron Lett.* 1988, 29, 4525) of **5b** was quenched with 40% aqueous methylamine instead of TEAB or H₂O, the γ -amidate **7b** was generated as the only detectable product. Similar reaction was reported recently for the preparation of the γ -amidate of pppA2'p5'A2'p5'A.¹²

Carboxylate group was introduced into 5-position of uridine both on the nucleoside level and post-synthetically (Method C) (Scheme 3). 5-Iodo-2'-deoxy-2'-fluorouridine (**16**) was coupled with methyl acrylate using modified Heck reaction¹³ to yield **17** in 85% yield. 5'-*O*-Dimethoxytritylation, followed by *in situ* 3'-*O*-acetylation and subsequent detritylation afforded 3'-protected derivative **18**. Phosphorylation using 2-chloro-4*H*-1,3,2-benzodioxaphosphorin-4-one followed by pyrophosphate addition and oxidation (Ludwig, J.; Eckstein, F. *J. Org. Chem.* 1989, 54, 631) afforded the desired triphosphate in 54% yield. On the other hand, 5-(3-aminopropyl)uridine 5'-triphosphate **6b** was coupled with *N*-hydroxysuccinimide ester of Fmoc-Asp-OFm to afford, after removal of Fmoc and Fm groups with diethylamine, the desired aminoacyl conjugate **20** in 50% yield.

Cytidine derivatives comprising 3-aminopropyl and 3(*N*-succinyl)aminopropyl groups were synthesized according to Scheme 4. Peracylated 5-(3-aminopropynyl)uracil derivative **2b** is reduced using catalytic hydrogenation and then converted in seven steps and 5% overall yield into 3'-acetylated cytidine derivative **25**. This synthesis was plagued by poor solubility of intermediates and formation of the N⁴-cyclized byproduct during ammonia treatment of the 4-triazolyl intermediate. Phosphorylation of **25** as described in reference 11 yielded triphosphate **26** and N⁴-cyclized product **27** in 1:1 ratio. They were easily separated on Sephadex DEAE A-25 ion exchange column using 0.1-0.8M TEAB gradient. It appears that under basic conditions the free primary amine can displace any remaining intact 4-NHBz group leading to the cyclized product. This is similar to displacement of 4-triazolyl group by primary amine as mentioned above.

We reasoned that utilization of *N*⁴-unprotected cytidine will solve this problem. This lead to an improved synthesis of 26: Iodination of 2'-deoxy-2'-fluorocytidine (28) provided the 5-iodo derivative 29 in 58% yield. This compound was then smoothly converted into 5-(3-aminopropynyl) derivative 30. Hydrogenation afforded 5-(3-aminopropyl) derivative 31 which was phosphorylated directly with POCl₃/ PPI to afford 26 in 37% yield. Coupling of the 5'-triphosphate 26 with succinic anhydride yielded succinylated derivative 32 in 36% yield.

Synthesis of 5-Imidazoleacetic acid 2'-deoxy-5'-triphosphate uridine

5-dintrophenylimidazoleacetic acid 2'-deoxy uridine nucleoside (80 mg) was dissolved in 5 ml of triethylphosphate while stirring under argon, and the reaction mixture was cooled to 0°C. Phosphorous oxychloride (1.8 eq, 22 ml) was added to the reaction mixture at 0°C, three more aliquots were added over the course of 48 hours at room temperature. The reaction mixture was then diluted with anhydrous MeCN (5 ml) and cooled to 0°C, followed by the addition of tributylamine (0.65 ml) and tributylammonium pyrophosphate (4.0 eq, 0.24 g). After 45 minutes, the reaction was quenched with 10 ml aq. methyl amine for four hours. After co-evaporation with MeOH (3x), purified material on DEAE Sephadex followed by RP chromatography to afford 15 mg of triphosphate.

Synthesis of 2'-(*N*-lysyl)-amino-2'-deoxy-cytidine Triphosphate

2'-(*N*-lysyl)-amino-2'-deoxy cytidine (0.180 g, 0.22 mmol) was dissolved in triethyl phosphate (2.00 ml) under Ar. The solution was cooled to 0 °C in an ice bath. Phosphorus oxychloride (99.999%, 3 eq., 0.0672 mL) was added to the solution and the reaction was stirred for two hours at 0 °C. Tributylammonium pyrophosphate (4 eq., 0.400 g) was dissolved in 3.42 mL of acetonitrile and tributylamine (0.165 mL). Acetonitrile (1 mL) was added to the monophosphate solution followed by the pyrophosphate solution which was added dropwise. The resulting solution was clear. The reaction was allowed to warm up to room temperature. After stirring for 45 minutes, methylamine (5 mL) was added and the reaction and stirred at room temperature for 2 hours. A biphasic mixture appeared (little beads at the bottom of the flask). TLC (7:1:2 iPrOH:NH₄OH:H₂O) showed the appearance of triphosphate material. The solution was concentrated, dissolved in water and loaded on a newly prepared DEAE Sephadex A-25 column. The column was washed with a gradient up to 0.6 M TEAB buffer and the product eluted off in fractions

90-95. The fractions were analyzed by ion exchange HPLC. Each fraction showed one triphosphate peak that eluted at ~4.000 minutes. The fractions were combined and pumped down from methanol to remove buffer salt to yield 15.7 mg of product.

Synthesis of 2'-deoxy-2'-(L-histidine)amino Cytidine Triphosphate

- 5 2'-[N-Fmoc, *Nimid* -dinitrophenyl-histidyl]amino-2'-cytidine (0.310 g, 4.04 mmol) was dissolved in triethyl phosphate (3 ml) under Ar. The solution was cooled to 0 °C. Phosphorus oxychloride (1.8 eq., 0.068 mL) was added to the solution and stored overnight in the freezer. The next morning TLC (10% MeOH in CH₂Cl₂) showed significant starting material, one more equivalent of POCl₃ was added. After two hours,
- 10 TLC still showed starting material. Tributylamine (0.303 mL) and Tributylammonium pyrophosphate (4 eq., 0.734 g) dissolved in 6.3 mL of acetonitrile (added dropwise) were added to the monophosphate solution. The reaction was allowed to warm up to room temperature. After stirring for 15 min, methylamine (10 mL) was added at room temperature and stirring continued for 2 hours. TLC (7:1:2 iPrOH:NH₄OH:H₂O) showed
- 15 the appearance of triphosphate material. The solution was concentrated, dissolved in water and loaded on a DEAE Sephadex A-25 column. The column was washed with a gradient up to 0.6 M TEAB buffer and the product eluted off in fractions 170-179. The fractions were analyzed by ion exchange HPLC. Each fraction showed one triphosphate peak that eluted at ~6.77 minutes. The fractions were combined and pumped down from methanol
- 20 to remove buffer salt to afford 17 mg of product.

Screening for Novel Enzymatic nucleic acid molecule Motifs Using Modified NTPs (Class I Motif)

- Our initial pool contained 3×10^{14} individual sequences of 2'-amino-dCTP/2'-amino-dUTP RNA. We optimized transcription conditions in order to increase the amount
- 25 of RNA product by inclusion of methanol and lithium chloride. 2'-amino-2'-deoxynucleotides do not interfere with the reverse transcription and amplification steps of selection and confer nuclease resistance. We designed the pool to have two binding arms complementary to the substrate, separated by the random 40 nucleotide region. The 16-mer substrate had two domains, 5 and 10 nucleotides long, that bind the pool, separated by
- 30 an unpaired guanosine. On the 5' end of the substrate was a biotin attached by a C18 linker. This enabled us to link the substrate to a NeutrAvidin™ resin in a column format.

The desired reaction would be cleavage at the unpaired G upon addition of magnesium cofactor followed by dissociation from the column due to instability of the 5 base pair helix. A detailed protocol follows:

Enzymatic nucleic acid molecule Pool Prep: The initial pool DNA was prepared by
5 converting the following template oligonucleotides into double-stranded DNA by filling in
with taq polymerase. (template=5'-ACC CTC ACT AAA GGC CGT (N)₄₀ GGT TGC
ACA CCT TTC-3'; primer 1=5'-CAC TTA GCA TTA ACC CTC ACT AAA GGC
CGT-3'; primer 2=5'-TAA TAC GAC TCA CTA TAG GAA AGG TGT GCA ACC-3'.)
All DNA oligonucleotides were synthesized by Operon technologies. Template oligos
10 were purified by denaturing PAGE and Sep-pak chromatography columns (Waters). RNA
substrate oligos were using standard solid phase chemistry and purified by denaturing
PAGE followed by ethanol precipitation. Substrates for *in vitro* cleavage assays were 5'-
end labeled with gamma-³²P-ATP and T4 polynucleotide kinase followed by denaturing
PAGE purification and ethanol precipitation.

15 5 nmole of template, 10 nmole of each primer and 250 U taq polymerase were
incubated in a 10 ml volume with 1X PCR buffer (10 mM tris-HCl (pH 8.3), 1.5 mM
MgCl₂, 50 mM KCl) and 0.2 mM each dNTP as follows: 94°C, 4 minutes; (94°C, 1 min;
42°C, 1 min; 72°C, 2 min) through four cycles; and then 72°C, for 10 minutes. The
product was analyzed on 2% Separide™ agarose gel for size and then was extracted twice
20 with buffered phenol, then chloroform-isoamyl alcohol, and ethanol precipitated. The
initial RNA pool was made by transcription of 500 pmole (3 x 10¹⁴ molecules) of this
DNA as follows. Template DNA was added to 40 mM tris-HCl (pH 8.0), 12 mM MgCl₂,
5 mM dithiothreitol (DTT), 1 mM spermidine, 0.002% triton X-100, 1 mM LiCl, 4%
PEG-8000, 10% methanol, 2 mM ATP, 2 mM GTP, 2 mM 2'-amino-dCTP, 2 mM 2'-
25 amino-dUTP, 5 U/ml inorganic pyrophosphatase, and 5 U/μl T7 RNA polymerase at room
temperature for a total volume of 1 ml. A separate reaction contained a trace amount of
alpha-³²P-GTP for detection. Transcriptions were incubated at 37°C for 2 hours followed
by addition of equal volume STOP buffer (94% formamide, 20 mM EDTA, 0.05%
bromophenol blue). The resulting RNA was purified by 6% denaturing PAGE gel, Sep-
30 pak™ chromatography, and ethanol precipitated.

INITIAL SELECTION: 2 nmole of 16 mer 5'-biotinylated substrate (5'-biotin-C18
linker-GCC GUG GGU UGC ACA C-3') was linked to 200 μl UltraLink Immobilized

NeutrAvidin™ resin (400 µl slurry, Pierce) in binding buffer (20 mM NaPO₄ (pH 7.5), 150 mM NaCl) for 30 minutes at room temperature. The resulting substrate column was washed with 2 ml binding buffer followed by 2 ml column buffer (50 mM tris-HCl (pH 8.5), 100 mM NaCl, 50 mM KCl). The flow was capped off and 1000 pmole of initial
5 pool RNA in 200 µl column buffer was added to the column and incubated 30 minutes at room temperature. The column was uncapped and washed with 2 ml column buffer, then capped off. 200 µl elution buffer (=column buffer + 25 mM MgCl₂) was added to the column and allowed to incubate 30 minutes at room temperature. The column was uncapped and eluent collected followed by three 200 µl elution buffer washes. The
10 eluent/washes were ethanol precipitated using glycogen as carrier and rehydrated in 50 µl sterile H₂O. The eluted RNA was amplified by standard reverse transcription/PCR amplification techniques. 5-31 µl RNA was incubated with 20 pmol of primer 1 in 14 µl volume 90° for 3 min then placed on ice for 1 minute. The following reagent were added (final concentrations noted): 1X PCR buffer, 1 mM each dNTP, 2 U/µl RNase Inhibitor,
15 10 U/µl SuperScript™ II reverse transcriptase. The reaction was incubated 42° for 1 hour followed by 95° for 5 min in order to inactivate the reverse transcriptase. The volume was then increased to 100 µl by adding water and reagents for PCR: 1X PCR buffer, 20 pmol primer 2, and 2.5 U taq DNA polymerase. The reaction was cycled in a Hybaid thermocycler: 94°, 4 min; (94°C, 30 sec; 54°C, 30 sec; 72°C, 1 min) X 25; 72°C, 5 min.
20 Products were analyzed on agarose gel for size and ethanol precipitated. One-third to one-fifth of the PCR DNA was used to transcribe the next generation, in 100 µl volume, as described above. Subsequent rounds used 20 pmol RNA for the column with 40 pmol substrate.

TWO COLUMN SELECTION: At generation 8 (G8), the column selection was
25 changed to the two column format. 200 pmoles of 22 mer 5'-biotinylated substrate (5'-biotin-C18 linker-GCC GUG GGU UGC ACA CCU UUC C-C18 linker-thiol modifier C6 S-S-inverted abasic-3') was used in the selection column as described above. Elution was in 200 µl elution buffer followed by a 1 ml elution buffer wash. The 1200 µl eluent was passed through a product trap column by gravity. The product trap column was prepared
30 as follows: 200 pmol 16 mer 5'-biotinylated "product" (5'-GGU UGC ACA CCU UUC C-C18 linker-biotin-3') was linked to the column as described above and the column was equilibrated in elution buffer. Eluent from the product column was precipitated as

previously described. The products were amplified as above only with 2.5-fold more volume and 100 pmol each primer. 100 μ l of the PCR reaction was used to do a cycle course; the remaining fraction was amplified the minimal number of cycles needed for product. After 3 rounds (G11), there was visible activity in a single turnover cleavage
5 assay. By generation 13, 45% of the substrate was cleaved at 4 hours; k_{obs} of the pool was 0.037 min^{-1} in 25 mM MgCl_2 . We subcloned and sequenced generation 13; the pool was still very diverse. Since our goal was a enzymatic nucleic acid molecule that would work in a physiological environment, we decided to change selection pressure rather than exhaustively catalog G13.

10 Reselection of the N40 pool was started from G12 DNA. Part of the G12 DNA was subjected to hypermutagenic PCR (Vartanian *et al.*, 1996, *Nucleic Acids Research* 24, 2627-2631) to introduce a 10% per position mutation frequency and was designated N40H. At round 19, part of the DNA was hypermutagenized again, giving N40M and N40HM (a total of 4 parallel pools). The column substrates remained the same; buffers
15 were changed and temperature of binding and elution was raised to 37°C. Column buffer was replaced by physiological buffer (50 mM tris-HCl (pH 7.5), 140 mM KCl, 10 mM NaCl) and elution buffer was replaced by 1 mM Mg buffer (physiological buffer + 1 mM MgCl_2). Amount of time allowed for the pool to bind the column was eventually reduced to 10 min and elution time was gradually reduced from 30 min to 20 sec. Between rounds
20 18 and 23, k_{obs} for the N40 pool stayed relatively constant at 0.035-0.04 min^{-1} . Generation 22 from each of the 4 pools was cloned and sequenced.

 CLONING AND SEQUENCING: Generations 13 and 22 were cloned using Novagen's Perfectly Blunt™ Cloning kit (pT7Blue-3 vector) following the kit protocol. Clones were screened for insert by PCR amplification using vector-specific primers.
25 Positive clones were sequenced using ABI Prism 7700 sequence detection system and vector-specific primer. Sequences were aligned using MacVector software; two-dimensional folding was performed using Mfold software (Zuker, 1989, *Science* 244, 48-52; Jaeger *et al.*, 1989, *Biochemistry* 86, 7706-7710; Jaeger *et al.*, 1989, R. F. Doolittle ed., *Methods in Enzymology*, 183, 281-306). Individual clone transcription units were
30 constructed by PCR amplification with 50 pmol each primer 1 and primer 2 in 1X PCR buffer, 0.2 mM each dNTP, and 2.5 U of taq polymerase in 100 μ l volume cycled as follows: 94°C, 4 min; (94°C, 30 sec; 54°C, 30 sec; 72°C, 1 min) X 20; 72°C, 5 min.

Transcription units were ethanol precipitated, rehydrated in 30 μ l H₂O, and 10 μ l was transcribed in 100 μ l volume and purified as previously described.

Thirty-six clones from each pool were sequenced and were found to be variations of the same consensus motif. Unique clones were assayed for activity in 1 mM MgCl₂ and physiological conditions; nine clones represented the consensus sequence and were used in subsequent experiments. There were no mutations that significantly increased activity; most of the mutations were in regions believed to be duplex, based on the proposed secondary structure. In order to make the motif shorter, we deleted the 3'-terminal 25 nucleotides necessary to bind the primer for amplification. The measured rates of the full length and truncated molecules were both 0.04 min⁻¹; thus we were able reduce the size of the motif from 86 to 61 nucleotides. The molecule was shortened even further by truncating base pairs in the stem loop structures as well as the substrate recognition arms to yield a 48 nucleotide molecule. In addition, many of the ribonucleotides were replaced with 2-*O*-methyl modified nucleotides to stabilize the molecule. An example of the new motif is given in Figure 13. Those of ordinary skill in the art will recognize that the molecule is not limited to the chemical modifications shown in the figure and that it represents only one possible chemically modified molecule.

Kinetic Analysis:

Single turnover kinetics were performed with trace amounts of 5'-³²P-labeled substrate and 10-1000 nM pool of enzymatic nucleic acid molecule. 2X substrate in 1X buffer and 2X pool/enzymatic nucleic acid molecule in 1X buffer were incubated separately 90° for 3 min followed by equilibration to 37° for 3 min. Equal volume of 2X substrate was added to pool/enzymatic nucleic acid molecule at t₀ and the reaction was incubated at 37°C. Time points were quenched in 1.2 vol STOP buffer on ice. Samples were heated to 90°C for 3 min prior to separation on 15% sequencing gels. Gels were imaged using a PhosphorImager and quantitated using ImageQuant™ software (Molecular Dynamics). Curves were fit to double-exponential decay in most cases, although some of the curves required linear fits.

STABILITY: Serum stability assays were performed as previously described (Beigelman *et al.*, 1995, *J. Biol. Chem.* 270, 25702-25708). 1 μ g of 5'-³²P-labeled

synthetic enzymatic nucleic acid molecule was added to 13 μ l cold and assayed for decay in human serum. Gels and quantitation were as described in kinetics section.

SUBSTRATE REQUIREMENTS: Table 60 outlines the substrate requirements for Class I motif. Substrates maintained Watson-Crick or wobble base pairing with mutant Class I constructs. Activity in single turnover kinetic assay is shown relative to wild type Class I and 22 mer substrate (50 mM Tris-HCL (pH 7.5), 140 mM KCl, 10 mM NaCl, 1 mM MgCl₂, 100 nM ribozyme, 5 nM substrate, 37°C).

RANDOM REGION MUTATION ALIGNMENT: Table 61 outlines the random region alignment of 134 clones from generation 22 (1.x = N40, 2.x = N40M, 3.x = N40H, 4.x = N40HM). The number of copies of each mutant is in parenthesis in the table, deviations from consensus are shown. Mutations that maintain base pair U19:A34 are shown in *italic*. Activity in single turnover kinetic assay is shown relative to the G22 pool rate (50 mM Tris-HCL pH 7.5, 140 mM KCl, 10 mM NaCl, 1 mM MgCl₂, 100 nM ribozyme, trace substrate, 37°C).

STEM TRUNCATION AND LOOP REPLACEMENT ANALYSIS: Figure 25 shows a representation of Class I ribozyme stem truncation and loop replacement analysis. The K_{rel} is compared to a 61 mer Class I ribozyme measured as described above. Figure 26 shows examples of Class I ribozymes with truncated stem(s) and/or non-nucleotide linker replaced loop structures.

20

Inhibition of HCV Using Class I (Amberzyme) Motif

During HCV infection, viral RNA is present as a potential target for enzymatic nucleic acid molecule cleavage at several processes: uncoating, translation, RNA replication and packaging. Target RNA may be more or less accessible to enzymatic nucleic acid molecule cleavage at any one of these steps. Although the association between the HCV initial ribosome entry site (IRES) and the translation apparatus is mimicked in the HCV 5'UTR/luciferase reporter system, these other viral processes are not represented in the OST7 system. The resulting RNA/protein complexes associated with the target viral RNA are also absent. Moreover, these processes may be coupled in an HCV-infected cell which could further impact target RNA accessibility. Therefore, we

30

tested whether enzymatic nucleic acid molecules designed to cleave the HCV 5'UTR could effect a replicating viral system.

Recently, Lu and Wimmer characterized an HCV-poliovirus chimera in which the poliovirus IRES was replaced by the IRES from HCV (Lu & Wimmer, 1996, *Proc. Natl. Acad. Sci. USA.* 93, 1412-1417). Poliovirus (PV) is a positive strand RNA virus like HCV, but unlike HCV is non-enveloped and replicates efficiently in cell culture. The HCV-PV chimera expresses a stable, small plaque phenotype relative to wild type PV.

The capability of the new enzymatic nucleic acid molecule motifs to inhibit HCV RNA intracellularly was tested using a dual reporter system that utilizes both firefly and Renilla luciferase (Figure 14). A number of enzymatic nucleic acid molecules having the new class I motif (Amberzyme) were designed and tested (Table 56). The Amberzyme ribozymes were targeted to the 5' HCV UTR region, which when cleaved, would prevent the translation of the transcript into luciferase. OST-7 cells were plated at 12,500 cells per well in black walled 96-well plates (Packard) in medium DMEM containing 10% fetal bovine serum, 1% pen/strep, and 1% L-glutamine and incubated at 37°C overnight. A plasmid containing T7 promoter expressing 5' HCV UTR and firefly luciferase (T7C1-341 (Wang *et al.*, 1993, *J. of Virol.* 67, 3338-3344)) was mixed with a pRLSV40 Renilla control plasmid (Promega Corporation) followed by enzymatic nucleic acid molecule, and cationic lipid to make a 5X concentration of the reagents (T7C1-341 (4 µg/ml), pRLSV40 renilla luciferase control (6 µg/ml), enzymatic nucleic acid molecule (250 nM), transfection reagent (28.5 µg/ml).

The complex mixture was incubated at 37°C for 20 minutes. The media was removed from the cells and 120 µl of Opti-mem media was added to the well followed by 30 µl of the 5X complex mixture. 150 µl of Opti-mem was added to the wells holding the untreated cells. The complex mixture was incubated on OST-7 cells for 4 hours, lysed with passive lysis buffer (Promega Corporation) and luminescent signals were quantified using the Dual Luciferase Assay Kit using the manufacturer's protocol (Promega Corporation). The data shown in Figure 15 is a dose curve of enzymatic nucleic acid molecule targeting site 146 of the HCV RNA and is presented as a ratio between the firefly and Renilla luciferase fluorescence. The enzymatic nucleic acid molecule was able to reduce the quantity of HCV RNA at all enzymatic nucleic acid molecule concentrations

yielding an IC_{50} of approximately 5 nM. Other sites were also efficacious (**Figure 16**), in particular enzymatic nucleic acid molecules targeting sites 133, 209, and 273 were also able to reduce HCV RNA compared to the irrelevant (IRR) controls.

5 Cleavage of Substrates Using Completely Modified class I (Amberzyme) enzymatic nucleic acid molecule

The ability of an enzymatic nucleic acid, which is modified at every 2' position to cleave a target RNA was tested to determine if any ribonucleotide positions are necessary in the Amberzyme motif. Enzymatic nucleic acid molecules were constructed with 2'-O-methyl, and 2'-amino (NH_2) nucleotides and included no ribonucleotides (**Table 56**; gene name: no ribo) and kinetic analysis was performed as described in example 13. 100 nM enzymatic nucleic acid was mixed with trace amounts of substrate in the presence of 1 mM $MgCl_2$ at physiological conditions (37°C). The Amberzyme with no ribonucleotide present in it has a K_{rel} of 0.13 compared to the enzymatic nucleic acid with a few
10 ribonucleotides present in the molecule shown in **Table 56 (ribo)**. This shows that
15 Amberzyme enzymatic nucleic acid molecule may not require the presence of 2'-OH groups within the molecule for activity.

Substrate Recognition Rules for Class II (zinzyme) enzymatic nucleic acid molecules

Class II (zinzyme) ribozymes were tested for their ability to cleave base-paired substrates with all sixteen possible combinations of bases immediately 5' and 3' proximal to the bulged cleavage site G. Ribozymes were identical in all remaining positions of their 7 base pair binding arms. Activity was assessed at two and twenty-four hour time points under standard reaction conditions [20 mM HEPES pH 7.4, 140 mM KCl, 10 mM NaCl, 1 mM MgCl₂, 1 mM CaCl₂ – 37⁰ C]. **Figure 19** shows the results of this study. Base paired substrate UGG (not shown in the figure) cleaved as poorly as CGG shown in the figure. The figure shows the cleavage site substrate triplet in the 5'- 3' direction and 2 and 24 hour time points are shown top to bottom respectively. The results indicate the cleavage site triplet is most active with a 5'- Y-G-H –3' (where Y is C or U and H is A, C or U with cleavage between G and H); however, activity is detected particularly with the 24 hour time point for most paired substrates. All positions outside of the cleavage triplet were found to tolerate any base pairings (data not shown).

All possible mispairs immediately 5' and 3' proximal to the bulged cleavage site G were tested to a class II ribozyme designed to cleave a 5'-C-G-C –3'. It was observed the 5' and 3' proximal sites are as active with G:U wobble pairs, in addition, the 5' proximal site will tolerate a mismatch with only a slight reduction in activity [data not shown].

Screening for Novel Enzymatic nucleic acid molecule Motifs (Class II Motifs)

The selections were initiated with pools of > 10¹⁴ modified RNA's of the following sequence: 5'-GGGAGGAGGAAGUGCCU (N)₃₅ UGCCGCGCUCGCUCCAGUCC-3'. The RNA was enzymatically generated using the mutant T7 Y639F RNA polymerase prepared by Rui Souza. The following modified NTP's were incorporated: 2'-deoxy-2'-fluoro-adenine triphosphate, 2'-deoxy-2'-fluoro-uridine triphosphate or 2'-deoxy-2'-fluoro-5-[(N-imidazole-4acetyl)propyl amine] uridine triphosphate, and 2'-deoxy-2'-amino-cytidine triphosphate; natural guanine triphosphate was used in all selections so that alpha –³²P-GTP could be used to label pool RNA's. RNA pools were purified by denaturing gel electrophoresis 8% polyacrilamide 7 M Urea.

The following target RNA (resin A) was synthesized and coupled to Iodoacetyl Ultralink™ resin (Pierce) by the supplier's procedure: 5' –b-L-GGACUGGGAGCGAGCGCGCGCAGGCACU GAAG-L-S-B-3'; where b is biotin (Glenn

Research cat# 10-1953-nn), L is polyethylene glycol spacer (Glenn Research cat# 10-1918-nn), S is thiol-modifier C6 S-S (Glenn Research cat# 10-1936-nn), B is a standard inverted deoxy abasic.

RNA pools were added to 100 μ l of 5 μ M Resin A in the buffer A (20 mM HEPES
5 pH 7.4, 140 mM KCL, 10 mM NaCl) and incubated at 22°C for 5 minutes. The
temperature was then raised to 37°C for 10 minutes. The resin was washed with 5 ml
buffer A. Reaction was triggered by the addition of buffer B(20 mM HEPES pH 7.4, 140
mM KCL, 10 mM NaCl, 1 mM MgCl₂, 1 mM CaCl₂). Incubation proceeded for 20
minutes in the first generation and was reduced progressively to 1 minute in the final
10 generations; with 13 total generations. The reaction eluent was collected in 5 M NaCl to
give a final concentration of 2 M NaCl. To this was added 100 μ l of 50% slurry Ultralink
NeutraAvidin™ (Pierce). Binding of cleaved biotin product to the avidin resin was
allowed by 20 minute incubation at 22°C. The resin was subsequently washed with 5 ml
of 20 mM HEPES pH 7.4, 2 M NaCl. Desired RNA's were removed by a 1.2 ml
15 denaturing wash 1M NaCl, 10 M Urea at 94°C over 10 minutes. RNA's were double
precipitated in 0.3 M sodium acetate to remove Cl⁻ ions inhibitory to reverse transcription.
Standard protocols of reverse transcription and PCR amplification were performed.
RNA's were again transcribed with the modified NTP's described above. After 13
generations cloning and sequencing provided 14 sequences which were able to cleave the
20 target substrate. Six sequences were characterized to determine secondary structure and
kinetic cleavage rates. The structures and kinetic data are given in Figure 17. The
sequences of eight other enzymatic nucleic acid molecule sequences are given in Table 57.
The size, sequence, and chemical compositions of these molecules can be modified as
described below or using other techniques well known in the art.

25

Nucleic Acid Catalyst Engineering

Sequence, chemical and structural variants of Class I and Class II enzymatic nucleic
acid molecule can be engineered and re-engineered using the techniques shown in this
application and known in the art. For example, the size of class I and class II enzymatic
30 nucleic acid molecules can, be reduced or increased using the techniques known in the art
(Zaug *et al.*, 1986, *Nature*, 324, 429; Ruffner *et al.*, 1990, *Biochem.*, 29, 10695; Beaudry *et*

al., 1990, *Biochem.*, 29, 6534; McCall *et al.*, 1992, *Proc. Natl. Acad. Sci., USA.*, 89, 5710; Long *et al.*, 1994, *supra*; Hendry *et al.*, 1994, *BBA* 1219, 405; Benseler *et al.*, 1993, *JACS*, 115, 8483; Thompson *et al.*, 1996, *Nucl. Acids Res.*, 24, 4401; Michels *et al.*, 1995, *Biochem.*, 34, 2965; Been *et al.*, 1992, *Biochem.*, 31, 11843; Guo *et al.*, 1995, *EMBO. J.*, 14, 368; Pan *et al.*, 1994, *Biochem.*, 33, 9561; Cech, 1992, *Curr. Op. Struc. Bio.*, 2, 605; Sugiyama *et al.*, 1996, *FEBS Lett.*, 392, 215; Beigelman *et al.*, 1994, *Bioorg. Med. Chem.*, 4, 1715; Santoro *et al.*, 1997, *PNAS* 94, 4262; all are incorporated in their totality by reference herein), to the extent that the overall catalytic activity of the ribozyme is not significantly decreased.

Further rounds of *in vitro* selection strategies described herein and variations thereof can be readily used by a person skilled in the art to evolve additional nucleic acid catalysts and such new catalysts are within the scope of the instant invention.

Example 16: Activity of Class II (zinzyme) nucleic acid catalysts to inhibit HER2 gene expression

Applicant has designed, synthesized and tested several class II (zinzyme) ribozymes targeted against HER2 RNA (see, for example, Tables 58, 59, and 62) in cell proliferation RNA reduction assays.

Proliferation assay:

The model proliferation assay used in the study can require a cell-plating density of 2000-10000 cells/well in 96-well plates and at least 2 cell doublings over a 5-day treatment period. Cells used in proliferation studies were either human breast or ovarian cancer cells (SKBR-3 and SKOV-3 cells respectively). To calculate cell density for proliferation assays, the FIPS (fluoro-imaging processing system) method well known in the art was used. This method allows for cell density measurements after nucleic acids are stained with CyQuant® dye, and has the advantage of accurately measuring cell densities over a very wide range 1,000-100,000 cells/well in 96-well format.

Ribozymes (50-200 nM) were delivered in the presence of cationic lipid at 2.0-5.0 µg/mL and inhibition of proliferation was determined on day 5 post-treatment. Two full ribozyme screens were completed resulting in the selection of 14 ribozymes. Class II (zinzyme) ribozymes against sites, 314 (RPI No. 18653), 443 (RPI No. 18680), 597 (RPI

No. 18697), 659 (RPI No. 18682), 878 (RPI Nos. 18683 and 18654), 881 (RPI Nos. 18684 and 18685) 934 (RPI No. 18651), 972 (RPI No. 18656, 19292, 19727, 19728, and 19293), 1292 (RPI No. 18726), 1541 (RPI No. 18687), 2116 (RPI No. 18729), 2932 (RPI No. 18678), 2540 (RPI No. 18715), and 3504 (RPI No. 18710) caused inhibition of proliferation ranging from 25-80% as compared to a scrambled control ribozyme. An example of results from a cell culture assay is shown in **Figure 20**. Referring to **Figure 20**, Class II ribozymes targeted against HER2 RNA are shown to cause significant inhibition of proliferation of cells. This shows that ribozymes, for instance the Class II (zinzyme) ribozymes are capable of inhibiting HER2 gene expression in mammalian cells.

RNA assay:

RNA was harvested 24 hours post-treatment using the Qiagen RNeasy® 96 procedure. Real time RT-PCR (TaqMan® assay) was performed on purified RNA samples using separate primer/probe sets specific for either target HER2 RNA or control actin RNA (to normalize for differences due to cell plating or sample recovery). Results are shown as the average of triplicate determinations of HER2 to actin RNA levels post-treatment. **Figure 30** shows class II ribozyme (zinzyme) mediated reduction in HER2 RNA targeting site 972 vs a scrambled attenuated control.

Dose response assays:

Active ribozyme was mixed with binding arm-attenuated control (BAC) ribozyme to a final oligonucleotide concentration of either 100, 200 or 400 nM and delivered to cells in the presence of cationic lipid at 5.0 µg/mL. Mixing active and BAC in this manner maintains the lipid to ribozyme charge ratio throughout the dose response curve. HER2 RNA reduction was measured 24 hours post-treatment and inhibition of proliferation was determined on day 5 post-treatment. The dose response antiproliferation results are summarized in **Figure 31** and the dose-dependent reduction of HER2 RNA results are summarized in **Figure 32**. **Figure 33** shows a combined dose response plot of both anti-proliferation and RNA reduction data for a class II ribozyme targeting site 972 of HER2 RNA (RPI 19293).

Example 17: Reduction of ribose residues in Class II (zinzyme) nucleic acid catalysts

Class II (zinzyme) nucleic acid catalysts were tested for their activity as a function of ribonucleotide content. A Zinzyme having no ribonucleotide residue (*ie.*, no 2'-OH group at the 2' position of the nucleotide sugar) against the K-Ras site 521 was designed. This 5 molecules were tested utilizing the chemistry shown in **Figure 27a**. The *in vitro* catalytic activity of the zinzyme construct was not significantly effected (the cleavage rate reduced only 10 fold).

The Kras zinzyme shown in **Figure 27a** was tested in physiological buffer with the divalent concentrations as indicated in the legend (high NaCl is an altered monovalent condition shown) of **Figure 28**. The 1 mM Ca^{++} condition yielded a rate of 0.005 min^{-1} while the 1 mM Mg^{++} condition yielded a rate of 0.002 min^{-1} . The ribose containing wild type yields a rate of 0.05 min^{-1} while substrate in the absence of zinzyme demonstrates less than 2% degradation at the longest time point under reaction conditions shown. This illustrates a well-behaved cleavage reaction done by a non-ribose containing catalyst with 15 only a 10-fold reduced cleavage as compared to ribonucleotide-containing zinzyme and vastly above non-catalyzed degradation.

A more detailed investigation into the role of ribose positions in the Class II (zinzyme) motif was carried out in the context of the HER2 site 972 (Applicant has further designed a fully modified Zinzyme as shown in **Figure 27b** targeting the HER2 RNA site 20 972). **Figure 29** is a diagram of the alternate formats tested and their relative rates of catalysis. The effect of substitution of ribose G for the 2'-O-methyl C-2'-O-methyl A in the loop of Zinzyme (see **Figure 34**) was insignificant when assayed with the Kras target but showed a modest rate enhancement in the HER2 assays. The activity of all Zinzyme motifs, including the fully stabilized "0 ribose" (RPI 19727) are well above background 25 noise level degradation. Zinzyme with only two ribose positions (RPI 19293) are sufficient to restore "wild-type" activity. Motifs containing 3 (RPI 19729), 4 (RPI 19730) or 5 ribose (RPI 19731) positions demonstrated a greater extent of cleavage and profiles almost identical to the 2 ribose motif. Applicant has thus demonstrated that a Zinzyme with no ribonucleotides present at any position can catalyze efficient RNA cleavage 30 activity. Thus, Zinzyme enzymatic nucleic acid molecules do not require the presence of 2'-OH group within the molecule for catalytic activity.

Example 18: Activity of reduced ribose containing Class II (zinzyme) nucleic acid catalysts to inhibit HER2 gene expression

A cell proliferation assay for testing reduced ribo class II (zinzyme) nucleic acid catalysts (50-400 nM) targeting HER2 site 972 was performed as described in example 19.

- 5 The results of this study are summarized in **Figure 35**. These results indicate significant inhibition of HER2 gene expression using stabilized Class II (zinzyme) motifs, including two ribo (RPI 19293), one ribo (RPI 19728), and non-ribo (RPI 19727) containing nucleic acid catalysts.

10 Applications

- The use of NTP's described in this invention have several research and commercial applications. These modified nucleotide triphosphates can be used for *in vitro* selection (evolution) of oligonucleotides with novel functions. Examples of *in vitro* selection protocols are incorporated herein by reference (Joyce, 1989, *Gene*, 82, 83-87; Beaudry *et al.*, 1992, *Science* 257, 635-641; Joyce, 1992, *Scientific American* 267, 90-97; Breaker *et al.*, 1994, *TIBTECH* 12, 268; Bartel *et al.*, 1993, *Science* 261:1411-1418; Szostak, 1993, *TIBS* 17, 89-93; Kumar *et al.*, 1995, *FASEB J.*, 9, 1183; Breaker, 1996, *Curr. Op. Biotech.*, 7, 442).

- 20 Additionally, these modified nucleotide triphosphates can be employed to generate modified oligonucleotide combinatorial chemistry libraries. Several references for this technology exist (Brenner *et al.*, 1992, *PNAS* 89, 5381-5383, Eaton, 1997, *Curr. Opin. Chem. Biol.* 1, 10-16), which are all incorporated herein by reference.

Diagnostic uses

- 25 Enzymatic nucleic acid molecules of this invention may be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of specific RNA in a cell. The close relationship between enzymatic nucleic acid molecule activity and the structure of the target RNA allows the detection of mutations in any region of the molecule which alters the base-pairing and three-dimensional structure of the target
30 RNA. By using multiple enzymatic nucleic acid molecules described in this invention, one may map nucleotide changes which are important to RNA structure and function *in vitro*, as well as in cells and tissues. Cleavage of target RNAs with enzymatic nucleic acid

molecules may be used to inhibit gene expression and define the role (essentially) of specified gene products in the progression of disease. In this manner, other genetic targets may be defined as important mediators of the disease. These experiments will lead to better treatment of the disease progression by affording the possibility of combinational therapies (e.g., multiple enzymatic nucleic acid molecules targeted to different genes, enzymatic nucleic acid molecules coupled with known small molecule inhibitors, radiation or intermittent treatment with combinations of enzymatic nucleic acid molecules and/or other chemical or biological molecules). Other *in vitro* uses of enzymatic nucleic acid molecules of this invention are well known in the art, and include detection of the presence of mRNAs associated with related conditions. Such RNA is detected by determining the presence of a cleavage product after treatment with a enzymatic nucleic acid molecule using standard methodology.

In a specific example, enzymatic nucleic acid molecules which can cleave only wild-type or mutant forms of the target RNA are used for the assay. The first enzymatic nucleic acid molecule is used to identify wild-type RNA present in the sample and the second enzymatic nucleic acid molecule will be used to identify mutant RNA in the sample. As reaction controls, synthetic substrates of both wild-type and mutant RNA will be cleaved by both enzymatic nucleic acid molecules to demonstrate the relative enzymatic nucleic acid molecule efficiencies in the reactions and the absence of cleavage of the "non-targeted" RNA species. The cleavage products from the synthetic substrates will also serve to generate size markers for the analysis of wild type and mutant RNAs in the sample population. Thus each analysis can involve two enzymatic nucleic acid molecules, two substrates and one unknown sample which can be combined into six reactions. The presence of cleavage products can be determined using an RNase protection assay so that full-length and cleavage fragments of each RNA can be analyzed in one lane of a polyacrylamide gel. It is not absolutely required to quantify the results to gain insight into the expression of mutant RNAs and putative risk of the desired phenotypic changes in target cells. The expression of mRNA whose protein product is implicated in the development of the phenotype is adequate to establish risk. If probes of comparable specific activity are used for both transcripts, then a qualitative comparison of RNA levels will be adequate and will decrease the cost of the initial diagnosis. Higher mutant form to

wild-type ratios will be correlated with higher risk whether RNA levels are compared qualitatively or quantitatively.

Additional Uses

5 Potential usefulness of sequence-specific enzymatic nucleic acid molecules of the instant invention can have many of the same applications for the study of RNA that DNA restriction endonucleases have for the study of DNA (Nathans *et al.*, 1975 *Ann. Rev. Biochem.* 44:273). For example, the pattern of restriction fragments can be used to establish sequence relationships between two related RNAs, and large RNAs could be
10 specifically cleaved to fragments of a size more useful for study. The ability to engineer sequence specificity of the enzymatic nucleic acid molecule is ideal for cleavage of RNAs of unknown sequence. Applicant describes the use of nucleic acid molecules to down-regulate gene expression of target genes in bacterial, microbial, fungal, viral, and eukaryotic systems including plant, or mammalian cells.

15 All patents and publications mentioned in the specification are indicative of the levels of skill of those skilled in the art to which the invention pertains. All references cited in this disclosure are incorporated by reference to the same extent as if each reference had been incorporated by reference in its entirety individually.

 One skilled in the art would readily appreciate that the present invention is well
20 adapted to carry out the objects and obtain the ends and advantages mentioned, as well as those inherent therein. The methods and compositions described herein as presently representative of preferred embodiments are exemplary and are not intended as limitations on the scope of the invention. Changes therein and other uses will occur to those skilled in the art, which are encompassed within the spirit of the invention, are defined by the scope
25 of the claims.

 It will be readily apparent to one skilled in the art that varying substitutions and modifications may be made to the invention disclosed herein without departing from the scope and spirit of the invention. Thus, such additional embodiments are within the scope of the present invention and the following claims.

30 The invention illustratively described herein suitably may be practiced in the absence of any element or elements, limitation or limitations which is not specifically disclosed herein. Thus, for example, in each instance herein any of the terms "comprising",

“consisting essentially of” and “consisting of” may be replaced with either of the other two terms. The terms and expressions which have been employed are used as terms of description and not of limitation, and there is no intention that in the use of such terms and expressions of excluding any equivalents of the features shown and described or portions thereof, but it is recognized that various modifications are possible within the scope of the invention claimed. Thus, it should be understood that although the present invention has been specifically disclosed by preferred embodiments, optional features, modification and variation of the concepts herein disclosed may be resorted to by those skilled in the art, and that such modifications and variations are considered to be within the scope of this invention as defined by the description and the appended claims.

In addition, where features or aspects of the invention are described in terms of Markush groups or other grouping of alternatives, those skilled in the art will recognize that the invention is also thereby described in terms of any individual member or subgroup of members of the Markush group or other group.

Thus, additional embodiments are within the scope of the invention and within the following claims

Table 1

TABLE 1**Characteristics of naturally occurring ribozymes****Group I Introns**

- Size: ~150 to >1000 nucleotides.
- Requires a U in the target sequence immediately 5' of the cleavage site.
- Binds 4-6 nucleotides at the 5'-side of the cleavage site.
- Reaction mechanism: attack by the 3'-OH of guanosine to generate cleavage products with 3'-OH and 5'-guanosine.
- Additional protein cofactors required in some cases to help folding and maintenance of the active structure.
- Over 300 known members of this class. Found as an intervening sequence in *Tetrahymena thermophila* rRNA, fungal mitochondria, chloroplasts, phage T4, blue-green algae, and others.
- Major structural features largely established through phylogenetic comparisons, mutagenesis, and biochemical studies [i,ii].
- Complete kinetic framework established for one ribozyme [iii,iv,v,vi].
- Studies of ribozyme folding and substrate docking underway [vii,viii,ix].
- Chemical modification investigation of important residues well established [x,xi].
- The small (4-6 nt) binding site may make this ribozyme too non-specific for targeted RNA cleavage, however, the *Tetrahymena* group I intron has been used to repair a "defective" β -galactosidase message by the ligation of new β -galactosidase sequences onto the defective message [xii].

RNAse P RNA (M1 RNA)

- Size: ~290 to 400 nucleotides.
- RNA portion of a ubiquitous ribonucleoprotein enzyme.
- Cleaves tRNA precursors to form mature tRNA [xiii].
- Reaction mechanism: possible attack by M^{2+} -OH to generate cleavage products with 3'-OH and 5'-phosphate.
- RNAse P is found throughout the prokaryotes and eukaryotes. The RNA subunit has been sequenced from bacteria, yeast, rodents, and primates.
- Recruitment of endogenous RNAse P for therapeutic applications is possible through hybridization of an External Guide Sequence (EGS) to the target RNA [xiv,xv]
- Important phosphate and 2' OH contacts recently identified [xvi,xvii]

Group II Introns

- Size: >1000 nucleotides.
- Trans cleavage of target RNAs recently demonstrated [xviii,xix].

Table 1

- Sequence requirements not fully determined.
- Reaction mechanism: 2'-OH of an internal adenosine generates cleavage products with 3'-OH and a "lariat" RNA containing a 3'-5' and a 2'-5' branch point.
- Only natural ribozyme with demonstrated participation in DNA cleavage [xx,xxi] in addition to RNA cleavage and ligation.
- Major structural features largely established through phylogenetic comparisons [xxii].
- Important 2' OH contacts beginning to be identified [xxiii]
- Kinetic framework under development [xxiv]

Neurospora VS RNA

- Size: ~144 nucleotides.
- Trans cleavage of hairpin target RNAs recently demonstrated [xxv].
- Sequence requirements not fully determined.
- Reaction mechanism: attack by 2'-OH 5' to the scissile bond to generate cleavage products with 2',3'-cyclic phosphate and 5'-OH ends.
- Binding sites and structural requirements not fully determined.
- Only 1 known member of this class. Found in *Neurospora* VS RNA.

Hammerhead Ribozyme

(see text for references)

- Size: ~13 to 40 nucleotides.
- Requires the target sequence UH immediately 5' of the cleavage site.
- Binds a variable number nucleotides on both sides of the cleavage site.
- Reaction mechanism: attack by 2'-OH 5' to the scissile bond to generate cleavage products with 2',3'-cyclic phosphate and 5'-OH ends.
- 14 known members of this class. Found in a number of plant pathogens (virusoids) that use RNA as the infectious agent.
- Essential structural features largely defined, including 2 crystal structures [xxvi,xxvii]
- Minimal ligation activity demonstrated (for engineering through *in vitro* selection) [xxviii]
- Complete kinetic framework established for two or more ribozymes [xxix].
- Chemical modification investigation of important residues well established [xxx].

Hairpin Ribozyme

- Size: ~50 nucleotides.
- Requires the target sequence GUC immediately 3' of the cleavage site.
- Binds 4-6 nucleotides at the 5'-side of the cleavage site and a variable number to the 3'-side of the cleavage site.
- Reaction mechanism: attack by 2'-OH 5' to the scissile bond to generate cleavage products with 2',3'-cyclic phosphate and 5'-OH ends.
- 3 known members of this class. Found in three plant pathogen (satellite RNAs of the tobacco ringspot virus, arabis mosaic virus and chicory yellow mottle virus) which uses RNA as the infectious agent.
- Essential structural features largely defined [xxxi,xxxii,xxxiii,xxxiv]

Table 1

- Ligation activity (in addition to cleavage activity) makes ribozyme amenable to engineering through *in vitro* selection [xxxv]
- Complete kinetic framework established for one ribozyme [xxxvi].
- Chemical modification investigation of important residues begun [xxxvii, xxxviii].

Hepatitis Delta Virus (HDV) Ribozyme

- Size: ~60 nucleotides.
- Trans cleavage of target RNAs demonstrated [xxxix].
- Binding sites and structural requirements not fully determined, although no sequences 5' of cleavage site are required. Folded ribozyme contains a pseudoknot structure [xl].
- Reaction mechanism: attack by 2'-OH 5' to the scissile bond to generate cleavage products with 2',3'-cyclic phosphate and 5'-OH ends.
- Only 2 known members of this class. Found in human HDV.
- Circular form of HDV is active and shows increased nuclease stability [xli]

-
- i. Michel, Francois; Westhof, Eric. Slippery substrates. *Nat. Struct. Biol.* (1994), 1(1), 5-7.
 - ii. Lisacek, Frederique; Diaz, Yolande; Michel, Francois. Automatic identification of group I intron cores in genomic DNA sequences. *J. Mol. Biol.* (1994), 235(4), 1206-17.
 - iii. Herschlag, Daniel; Cech, Thomas R.. Catalysis of RNA cleavage by the *Tetrahymena thermophila* ribozyme. 1. Kinetic description of the reaction of an RNA substrate complementary to the active site. *Biochemistry* (1990), 29(44), 10159-71.
 - iv. Herschlag, Daniel; Cech, Thomas R.. Catalysis of RNA cleavage by the *Tetrahymena thermophila* ribozyme. 2. Kinetic description of the reaction of an RNA substrate that forms a mismatch at the active site. *Biochemistry* (1990), 29(44), 10172-80.
 - v. Knitt, Deborah S.; Herschlag, Daniel. pH Dependencies of the *Tetrahymena* Ribozyme Reveal an Unconventional Origin of an Apparent pKa. *Biochemistry* (1996), 35(5), 1560-70.
 - vi. Bevilacqua, Philip C.; Sugimoto, Naoki; Turner, Douglas H.. A mechanistic framework for the second step of splicing catalyzed by the *Tetrahymena* ribozyme. *Biochemistry* (1996), 35(2), 648-58.
 - vii. Li, Yi; Bevilacqua, Philip C.; Mathews, David; Turner, Douglas H.. Thermodynamic and activation parameters for binding of a pyrene-labeled substrate by the *Tetrahymena* ribozyme: docking is not diffusion-controlled and is driven by a favorable entropy change. *Biochemistry* (1995), 34(44), 14394-9.
 - viii. Banerjee, Alope Raj; Turner, Douglas H.. The time dependence of chemical modification reveals slow steps in the folding of a group I ribozyme. *Biochemistry* (1995), 34(19), 6504-12.
 - ix. Zarrinkar, Patrick P.; Williamson, James R.. The P9.1-P9.2 peripheral extension helps guide folding of the *Tetrahymena* ribozyme. *Nucleic Acids Res.* (1996), 24(5), 854-8.
 - x. Strobel, Scott A.; Cech, Thomas R.. Minor groove recognition of the conserved G.cntdot.U pair at the *Tetrahymena* ribozyme reaction site. *Science* (Washington, D. C.) (1995), 267(5198), 675-9.
 - xi. Strobel, Scott A.; Cech, Thomas R.. Exocyclic Amine of the Conserved G.cntdot.U Pair at the Cleavage Site of the *Tetrahymena* Ribozyme Contributes to 5'-Splice Site Selection and Transition State Stabilization. *Biochemistry* (1996), 35(4), 1201-11.
 - xii. Sullenger, Bruce A.; Cech, Thomas R.. Ribozyme-mediated repair of defective mRNA by targeted trans-splicing. *Nature* (London) (1994), 371(6498), 619-22.
 - xiii. Robertson, H.D.; Altman, S.; Smith, J.D. *J. Biol. Chem.*, 247, 5243-5251 (1972).
 - xiv. Forster, Anthony C.; Altman, Sidney. External guide sequences for an RNA enzyme. *Science* (Washington, D. C., 1883-) (1990), 249(4970), 783-6.
 - xv. Yuan, Y.; Hwang, E. S.; Altman, S. Targeted cleavage of mRNA by human RNase P. *Proc. Natl. Acad. Sci. USA* (1992) 89, 8006-10.
 - xvi. Harris, Michael E.; Pace, Norman R.. Identification of phosphates involved in catalysis by the

Table 1

-
- ribozyme RNase P RNA. *RNA* (1995), 1(2), 210-18.
- xxv . Pan, Tao; Loria, Andrew; Zhong, Kun. Probing of tertiary interactions in RNA: 2'-hydroxyl-base contacts between the RNase P RNA and pre-tRNA. *Proc. Natl. Acad. Sci. U. S. A.* (1995), 92(26), 12510-14.
- xxviii . Pyle, Anna Marie; Green, Justin B.. Building a Kinetic Framework for Group II Intron Ribozyme Activity: Quantitation of Interdomain Binding and Reaction Rate. *Biochemistry* (1994), 33(9), 2716-25.
- xix . Michels, William J. Jr.; Pyle, Anna Marie. Conversion of a Group II Intron into a New Multiple-Turnover Ribozyme that Selectively Cleaves Oligonucleotides: Elucidation of Reaction Mechanism and Structure/Function Relationships. *Biochemistry* (1995), 34(9), 2965-77.
- xx . Zimmerly, Steven; Guo, Huatao; Eskes, Robert; Yang, Jian; Perlman, Philip S.; Lambowitz, Alan M.. A group II intron RNA is a catalytic component of a DNA endonuclease involved in intron mobility. *Cell* (Cambridge, Mass.) (1995), 83(4), 529-38.
- xxi . Griffin, Edmund A., Jr.; Qin, Zhifeng; Michels, Williams J., Jr.; Pyle, Anna Marie. Group II intron ribozymes that cleave DNA and RNA linkages with similar efficiency, and lack contacts with substrate 2'-hydroxyl groups. *Chem. Biol.* (1995), 2(11), 761-70.
- xxii . Michel, Francois; Ferat, Jean Luc. Structure and activities of group II introns. *Annu. Rev. Biochem.* (1995), 64, 435-61.
- xxiii . Abramovitz, Dana L.; Friedman, Richard A.; Pyle, Anna Marie. Catalytic role of 2'-hydroxyl groups within a group II intron active site. *Science* (Washington, D. C.) (1996), 271(5254), 1410-13.
- xxiv . Daniels, Danette L.; Michels, William J., Jr.; Pyle, Anna Marie. Two competing pathways for self-splicing by group II introns: a quantitative analysis of in vitro reaction rates and products. *J. Mol. Biol.* (1996), 256(1), 31-49.
- xxv . Guo, Hans C. T.; Collins, Richard A.. Efficient trans-cleavage of a stem-loop RNA substrate by a ribozyme derived from *Neurospora* VS RNA. *EMBO J.* (1995), 14(2), 368-76.
- xxvi . Scott, W.G., Finch, J.T., Aaron, K. The crystal structure of an all RNA hammerhead ribozyme: A proposed mechanism for RNA catalytic cleavage. *Cell*, (1995), 81, 991-1002.
- xxvii . McKay, Structure and function of the hammerhead ribozyme: an unfinished story. *RNA*, (1996), 2, 395-403.
- xxviii . Long, D., Uhlenbeck, O., Hertel, K. Ligation with hammerhead ribozymes. US Patent No. 5,633,133.
- xxix . Hertel, K.J., Herschlag, D., Uhlenbeck, O. A kinetic and thermodynamic framework for the hammerhead ribozyme reaction. *Biochemistry*, (1994) 33, 3374-3385. Beigelman, L., *et al.*, Chemical modifications of hammerhead ribozymes. *J. Biol. Chem.*, (1995) 270, 25702-25708.
- xxx . Beigelman, L., *et al.*, Chemical modifications of hammerhead ribozymes. *J. Biol. Chem.*, (1995) 270, 25702-25708.
- xxxi . Hampel, Arnold; Tritz, Richard; Hicks, Margaret; Cruz, Phillip. 'Hairpin' catalytic RNA model: evidence for helices and sequence requirement for substrate RNA. *Nucleic Acids Res.* (1990), 18(2), 299-304.
- xxxii . Chowrira, Bharat M.; Berzal-Herranz, Alfredo; Burke, John M.. Novel guanosine requirement for catalysis by the hairpin ribozyme. *Nature* (London) (1991), 354(6351), 320-2.
- xxxiii . Berzal-Herranz, Alfredo; Joseph, Simpson; Chowrira, Bharat M.; Butcher, Samuel E.; Burke, John M.. Essential nucleotide sequences and secondary structure elements of the hairpin ribozyme. *EMBO J.* (1993), 12(6), 2567-73.
- xxxiv . Joseph, Simpson; Berzal-Herranz, Alfredo; Chowrira, Bharat M.; Butcher, Samuel E.. Substrate selection rules for the hairpin ribozyme determined by in vitro selection, mutation, and analysis of mismatched substrates. *Genes Dev.* (1993), 7(1), 130-8.
- xxxv . Berzal-Herranz, Alfredo; Joseph, Simpson; Burke, John M.. In vitro selection of active hairpin ribozymes by sequential RNA-catalyzed cleavage and ligation reactions. *Genes Dev.* (1992), 6(1), 129-34.
- xxxvi . Hegg, Lisa A.; Fedor, Martha J.. Kinetics and Thermodynamics of Intermolecular Catalysis by Hairpin Ribozymes. *Biochemistry* (1995), 34(48), 15813-28.
- xxxvii . Grasby, Jane A.; Mersmann, Karin; Singh, Mohinder; Gait, Michael J.. Purine Functional Groups in Essential Residues of the Hairpin Ribozyme Required for Catalytic Cleavage of RNA. *Biochemistry*

Table 1

(1995), 34(12), 4068-76.

^{xxxviii} . Schmidt, Sabine; Beigelman, Leonid; Karpeisky, Alexander; Usman, Nassim; Sorensen, Ulrik S.; Gait, Michael J.. Base and sugar requirements for RNA cleavage of essential nucleoside residues in internal loop B of the hairpin ribozyme: implications for secondary structure. *Nucleic Acids Res.* (1996), 24(4), 573-81.

^{xxxix} . Perrotta, Anne T.; Been, Michael D.. Cleavage of oligoribonucleotides by a ribozyme derived from the hepatitis delta virus RNA sequence. *Biochemistry* (1992), 31(1), 16-21.

^{xl} . Perrotta, Anne T.; Been, Michael D.. A pseudoknot-like structure required for efficient self-cleavage of hepatitis delta virus RNA. *Nature (London)* (1991), 350(6317), 434-6.

^{xli} . Puttaraju, M.; Perrotta, Anne T.; Been, Michael D.. A circular trans-acting hepatitis delta virus ribozyme. *Nucleic Acids Res.* (1993), 21(18), 4253-8.

Table 2

Table 2:

A. 2.5 μ mol Synthesis Cycle ABI 394 Instrument

Reagent	Equivalents	Amount	Wait Time* DNA	Wait Time* 2'-O-methyl	Wait Time* RNA
Phosphoramidites	6.5	163 μ L	45 sec	2.5 min	7.5 min
S-Ethyl Tetrazole	23.8	238 μ L	45 sec	2.5 min	7.5 min
Acetic Anhydride	100	233 μ L	5 sec	5 sec	5 sec
N-Methyl Imidazole	186	233 μ L	5 sec	5 sec	5 sec
TCA	176	2.3 mL	21 sec	21 sec	21 sec
Iodine	11.2	1.7 mL	45 sec	45 sec	45 sec
Beaucage	12.9	645 μ L	100 sec	300 sec	300 sec
Acetonitrile	NA	6.67 mL	NA	NA	NA

B. 0.2 μ mol Synthesis Cycle ABI 394 Instrument

Reagent	Equivalents	Amount	Wait Time* DNA	Wait Time* 2'-O-methyl	Wait Time* RNA
Phosphoramidites	15	31 μ L	45 sec	233 sec	465 sec
S-Ethyl Tetrazole	38.7	31 μ L	45 sec	233 min	465 sec
Acetic Anhydride	655	124 μ L	5 sec	5 sec	5 sec
N-Methyl Imidazole	1245	124 μ L	5 sec	5 sec	5 sec
TCA	700	732 μ L	10 sec	10 sec	10 sec
Iodine	20.6	244 μ L	15 sec	15 sec	15 sec
Beaucage	7.7	232 μ L	100 sec	300 sec	300 sec

Table 2

C. 0.2 μ mol Synthesis Cycle 96 well Instrument						
Reagent	Equivalents DNA/2'-O-methyl/Ribo	Amount DNA/2'-O-methyl/Ribo	Wait Time* DNA	Wait Time* 2'-O- methyl	Wait Time* Ribo	
Acetonitrile	NA	2.64 mL	NA	NA	NA	
Phosphoramidites	22/33/66	40/60/120 μ L	60 sec	180 sec	360sec	
S-Ethyl Tetrazole	70/105/210	40/60/120 μ L	60 sec	180 min	360 sec	
Acetic Anhydride	265/265/265	50/50/50 μ L	10 sec	10 sec	10 sec	
N-Methyl Imidazole	502/502/502	50/50/50 μ L	10 sec	10 sec	10 sec	
TCA	238/475/475	250/500/500 μ L	15 sec	15 sec	15 sec	
Iodine	6.8/6.8/6.8	80/80/80 μ L	30 sec	30 sec	30 sec	
Beaucage	34/51/51	80/120/120	100 sec	200 sec	200 sec	
Acetonitrile	NA	1150/1150/1150 μ L	NA	NA	NA	

* Wait time does not include contact time during delivery.

Table 3

Table 3: Human PTP-1B Hammerhead Ribozyme and Target Sequence

Nt. Position	Ribozyme Sequence	Seq. ID Nos.	Substrate Sequence	Seq. ID Nos.
15	UGCCGCUC CUGAUGAG X CGAA AGGCCGCG	1	CGCGGCCT A GAGCGGCA	529
72	AUCUCCAU CUGAUGAG X CGAA ACGGGCCA	2	TGGCCCGT C ATGGAGAT	530
92	UCUGCUCG CUGAUGAG X CGAA ACUCCUUU	3	AAAGGAGT T CGAGCAGA	531
93	AUCUGCUC CUGAUGAG X CGAA AACUCCUU	4	AAGGAGTT C GAGCAGAT	532
102	GACUUGUC CUGAUGAG X CGAA AUCUGCUC	5	GAGCAGAT C GACAAGTC	533
110	AGCUCCCG CUGAUGAG X CGAA ACUUGUCG	6	CGACAAGT C CGGGAGCT	534
129	UCCUGGUA CUGAUGAG X CGAA AUGGCCGC	7	GCGGCCAT T TACCAGGA	535
130	AUCCUGGU CUGAUGAG X CGAA AAUGGCCG	8	CGGCCATT T ACCAGGAT	536
131	UAUCCUGG CUGAUGAG X CGAA AAAUGGCC	9	GGCCATTT A CCAGGATA	537
139	AUGUCGGA CUGAUGAG X CGAA AUCCUGGU	10	ACCAGGAT A TCCGACAT	538
141	UCAUGUCG CUGAUGAG X CGAA AUAUCCUG	11	CAGGATAT C GCACATGA	539
161	UACAUGGG CUGAUGAG X CGAA AGUCACUG	12	CAGTGACT T CCCATGTA	540
162	CUACAUGG CUGAUGAG X CGAA AAGUCACU	13	AGTGACTT C CCATGTAG	541
169	GGCCACUC CUGAUGAG X CGAA ACAUGGGA	14	TCCCATGT A GAGTGGCC	542
183	UUCUUAGG CUGAUGAG X CGAA AGCUUGGC	15	GCCAAGCT T CCTAAGAA	543
184	GUUCUAG CUGAUGAG X CGAA AAGCUUGG	16	CCAAGCTT C CTAAGAAC	544
187	UUUGUUCU CUGAUGAG X CGAA AGGAAGCU	17	AGCTTCCT A AGAACAAA	545
205	UCUGUACC CUGAUGAG X CGAA AUUUCGGU	18	ACCGAAT A GGTACAGA	546
209	CGUCUCUG CUGAUGAG X CGAA ACCUAUUU	19	AAATAGGT A CAGAGACG	547
219	AAGGGACU CUGAUGAG X CGAA ACGUCUCU	20	AGAGACGT C AGTCCCTT	548
223	GUCAAAGG CUGAUGAG X CGAA ACUGACGU	21	ACGTCAGT C CCTTTGAC	549
227	UAUGGUCA CUGAUGAG X CGAA AGGGACUG	22	CAGTCCCT T TGACCATA	550
228	CUAUGGUC CUGAUGAG X CGAA AAGGGACU	23	AGTCCCTT T GACCATAG	551
235	AAUCCGAC CUGAUGAG X CGAA AUGGUCAA	24	TTGACCAT A GTCGGATT	552
238	UUUAAUCC CUGAUGAG X CGAA ACUAUGGU	25	ACCATAGT C GGATTAAA	553
243	UGUAGUUU CUGAUGAG X CGAA AUCCGACU	26	AGTCGGAT T AAATAACA	554
244	AUGUAGUU CUGAUGAG X CGAA AAUCCGAC	27	GTCGGATT A AACTACAT	555
249	UCUGAUG CUGAUGAG X CGAA AGUUUAAU	28	ATTAAACT A CATCAAGA	556
253	AUCUUCUU CUGAUGAG X CGAA AUGUAGUU	29	AACTACAT C AAGAAGAT	557
262	AUAGUCAU CUGAUGAG X CGAA AUCUUCUU	30	AAGAAGAT A ATGACTAT	558
269	CGUUGAUA CUGAUGAG X CGAA AGUCAUUA	31	TAATGACT A TATCAACG	559
271	AGCGUUGA CUGAUGAG X CGAA AUAGUCAU	32	ATGACTAT A TCAACGCT	560
273	CUAGCGUU CUGAUGAG X CGAA AUAUAGUC	33	GACTATAT C AACGCTAG	561
280	UAUCAAAC CUGAUGAG X CGAA AGCGUUGA	34	TCAACGCT A GTTTGATA	562
283	UUUUAUCA CUGAUGAG X CGAA ACUAGCGU	35	ACGCTAGT T TGATAAAA	563
284	UUUUUAUC CUGAUGAG X CGAA AACUAGCG	36	CGCTAGTT T GATAAAAA	564
288	UCCAUUUU CUGAUGAG X CGAA AUCAAACU	37	AGTTTGAT A AAAATGGA	565
313	AAGAAUGU CUGAUGAG X CGAA ACUCUUUU	38	AAAGGAGT T ACATTCTT	566
314	UAAGAAUG CUGAUGAG X CGAA AACUCCUU	39	AAGGAGTT A CATTCTTA	567
318	UGGGUAAAG CUGAUGAG X CGAA AUGUAAACU	40	AGTTACAT T CTTACCCA	568
319	CUGGGUAA CUGAUGAG X CGAA AAUGUAAAC	41	GTTACATT C TTACCCAG	569
321	CCCUUGGU CUGAUGAG X CGAA AGAAUGUA	42	TACATTCT T ACCAGGG	570
322	GCCUUGGG CUGAUGAG X CGAA AAGAAUGU	43	ACATTCTT A CCCAGGGC	571
334	GUUAGGCA CUGAUGAG X CGAA AGGGCCCU	44	AGGGCCCT T TGCCTAAC	572

Table 3

335	UGUUAGGC CUGAUGAG X CGAA AAGGGCCC	45	GGGCCCTT T GCCTAACA	573
340	GCAUGUGU CUGAUGAG X CGAA AGGCAAAG	46	CTTTGCCT A ACACATGC	574
352	CCAAAAGU CUGAUGAG X CGAA ACCGCAUG	47	CATGCGGT C ACTTTTGG	575
356	UCUCCCAA CUGAUGAG X CGAA AGUGACCG	48	CGGTCACT T TTGGGAGA	576
357	AUCUCCCA CUGAUGAG X CGAA AAGUGACC	49	GGTCACTT T TGGGAGAT	577
358	CAUCUCCC CUGAUGAG X CGAA AAAGUGAC	50	GTCACCTT T GGGAGATG	578
393	AGCAUGAC CUGAUGAG X CGAA ACACCCCU	51	AGGGGTGT C GTCATGCT	579
396	UUGAGCAU CUGAUGAG X CGAA ACGACACC	52	GGTGTGCT C ATGCTCAA	580
402	ACUCUGUU CUGAUGAG X CGAA AGCAUGAC	53	GTCATGCT C AACAGAGT	581
424	UUUUAACG CUGAUGAG X CGAA ACCUUUCU	54	AGAAAGGT T CGTTAAAA	582
425	AUUUUAAC CUGAUGAG X CGAA AACCUIUC	55	GAAAGGTT C GTTAAAT	583
428	CGCAUUUU CUGAUGAG X CGAA ACGAACCU	56	AGGTTCGT T AAAATGCG	584
429	GCGCAUUU CUGAUGAG X CGAA AACGAACC	57	GGTTCGTT A AAATGCGC	585
443	GUGGCCAG CUGAUGAG X CGAA AUUGUGCG	58	CGCACAAT A CTGGCCAC	586
474	UCUUCAAA CUGAUGAG X CGAA AUCAUCUC	59	GAGATGAT C TTTGAAGA	587
476	UGUCUUCA CUGAUGAG X CGAA AGAUCauc	60	GATGATCT T TGAAGACA	588
477	GUGUCUUC CUGAUGAG X CGAA AAGAUCAU	61	ATGATCTT T GAAGACAC	589
490	UAAUUUCA CUGAUGAG X CGAA AUUUGUGU	62	ACACAAAT T TGAAATTA	590
491	UUAUUUUC CUGAUGAG X CGAA AAUUGUG	63	CACAAATT T GAAATTAA	591
497	UCAAUGUU CUGAUGAG X CGAA AUUUCAAA	64	TTTGAAAT T AACATTGA	592
498	AUCAUGU CUGAUGAG X CGAA AAUUCUAA	65	TTGAAATT A ACATTGAT	593
503	CAGAGAUC CUGAUGAG X CGAA AUGUUAU	66	ATTAACAT T GATCTCTG	594
507	UCUUCAGA CUGAUGAG X CGAA AUCAAUGU	67	ACATTGAT C TCTGAAGA	595
509	UAUCUUCA CUGAUGAG X CGAA AGAUCUUA	68	ATTGATCT C TGAAGATA	596
517	UGACUUGA CUGAUGAG X CGAA AUCUUCAG	69	CTGAAGAT A TCAAGTCA	597
519	UAUGACUU CUGAUGAG X CGAA AUAUCUUC	70	GAAGATAT C AAGTCATA	598
524	UAUAUAU CUGAUGAG X CGAA ACUUGAUA	71	TATCAAGT C ATATTATA	599
527	CUGUAUAA CUGAUGAG X CGAA AUGACUUG	72	CAAGTCAT A TTATACAG	600
529	CACUGUAU CUGAUGAG X CGAA AUAUGACU	73	AGTCATAT T ATACAGTG	601
530	GCACUGUA CUGAUGAG X CGAA AAUAUGAC	74	GTCATATT A TACAGTGC	602
532	UCGCACUG CUGAUGAG X CGAA AUAUAUG	75	CATATTAT A CAGTGCGA	603
546	UCCAAUUC CUGAUGAG X CGAA AGCUGUCG	76	CGACAGCT A GAATTGGA	604
551	GGUUUUC CUGAUGAG X CGAA AUUCUAGC	77	GCTAGAAT T GGAAAACC	605
561	UGGGUUGU CUGAUGAG X CGAA AGGUUUUC	78	GAAAACCT T ACAACCCA	606
562	UUGGUUG CUGAUGAG X CGAA AAGGUUUU	79	AAAACCTT A CAACCCAA	607
577	GAUCUCUC CUGAUGAG X CGAA AGUUUCUU	80	AAGAACT C GAGAGATC	608
585	AAAUGUAA CUGAUGAG X CGAA AUCUCUCG	81	CGAGAGAT C TTACATTT	609
587	GGAAUGU CUGAUGAG X CGAA AGAUCUCU	82	AGAGATCT T ACATTTCC	610
588	UGGAAUG CUGAUGAG X CGAA AAGAUUC	83	GAGATCTT A CATTTCCA	611
592	AUAGUGGA CUGAUGAG X CGAA AUGUAAGA	84	TCTTACAT T TCCACTAT	612
593	UAUAGUGG CUGAUGAG X CGAA AAUGUAAG	85	CTTACATT T CCACTATA	613
594	GUAUAGUG CUGAUGAG X CGAA AAAUGUAA	86	TTACATT C CACTATAC	614
599	AUGUGGUA CUGAUGAG X CGAA AGUGGAAA	87	TTTCCACT A TACCACAT	615
601	CCAUGUGG CUGAUGAG X CGAA AUAGUGGA	88	TCCACTAT A CCACATGG	616
617	GGACUCCA CUGAUGAG X CGAA AGUCAGGC	89	GCCTGACT T TGGAGTCC	617
618	GGGACUCC CUGAUGAG X CGAA AAGUCAGG	90	CCTGACTT T GGAGTCCC	618
624	GAUUCAGG CUGAUGAG X CGAA ACUCCAAA	91	TTTGAGT C CCTGAATC	619

Table 3

632	AGGCUGGU CUGAUGAG X CGAA AUUCAGGG	92	CCCTGAAT C ACCAGCCT	620
641	UCAAGAAU CUGAUGAG X CGAA AGGCUGGU	93	ACCAGCCT C ATTCTTGA	621
644	AGUUCAAG CUGAUGAG X CGAA AUGAGGCU	94	AGCCTCAT T CTGAACT	622
645	AAGUUCAA CUGAUGAG X CGAA AAUGAGGC	95	GCCTCATT C TTGAACTT	623
647	GAAAGUUC CUGAUGAG X CGAA AGAAUGAG	96	CTCATTCT T GAACTTTC	624
653	UGAAAAGA CUGAUGAG X CGAA AGUUCAAG	97	CTTGAAC T TCTTTTCA	625
654	UUGAAAAG CUGAUGAG X CGAA AAGUUCAA	98	TTGAACTT T CTTTTC	626
655	UUUGAAAA CUGAUGAG X CGAA AAAGUUCA	99	TGAACTTT C TTTTCAAA	627
657	ACUUUGAA CUGAUGAG X CGAA AGAAAGUU	100	AACTTTCT T TTCAAAGT	628
658	GACUUUGA CUGAUGAG X CGAA AAGAAAGU	101	ACTTTCTT T TCAAAGTC	629
659	GGACUUUG CUGAUGAG X CGAA AAAGAAAG	102	CTTTCTTT T CAAAGTCC	630
660	CGGACUUU CUGAUGAG X CGAA AAAAGAAA	103	TTTCTTTT C AAAGTCCG	631
666	GACUCUCG CUGAUGAG X CGAA ACUUUGAA	104	TTCAAAGT C CGAGAGTC	632
674	GUGACCCU CUGAUGAG X CGAA ACUCUCGG	105	CCGAGAGT C AGGGTCAC	633
680	GGCUGAGU CUGAUGAG X CGAA ACCCUGAC	106	GTCAGGGT C ACTCAGCC	634
684	UCCGGGCU CUGAUGAG X CGAA AGUGACCC	107	GGGTCACT C AGCCCGGA	635
705	UGCACCAC CUGAUGAG X CGAA ACGGGCCC	108	GGGCCCCG T GTGGTGCA	636
729	GACCUGCC CUGAUGAG X CGAA AUGCCUGC	109	GCAGGCAT C GGCAGGTC	637
737	AGGUUCCA CUGAUGAG X CGAA ACCUGCCG	110	CGGCAGGT C TGGAACCT	638
746	CCAGACAG CUGAUGAG X CGAA AGGUUCCA	111	TGGAACCT T CTGTCTGG	639
747	GCCAGACA CUGAUGAG X CGAA AAGGUUCC	112	GGAACCTT C TGTCTGGC	640
751	AUCAGCCA CUGAUGAG X CGAA ACAGAAGG	113	CCTTCTGT C TGCTGAT	641
760	GAGGCAGG CUGAUGAG X CGAA AUCAGCCA	114	TGGCTGAT A CCTGCCTC	642
768	AUCAGCAA CUGAUGAG X CGAA AGGCAGGU	115	ACCTGCCT C TTGCTGAT	643
770	CCAUCAGC CUGAUGAG X CGAA AGAGGCAG	116	CTGCCTCT T GCTGATGG	644
796	AACGGAAG CUGAUGAG X CGAA AGGGUCUU	117	AAGACCCT T CTTCGTT	645
797	CAACGGAA CUGAUGAG X CGAA AAGGGUCU	118	AGACCCTT C TTCCGTTG	646
799	AUCAACGG CUGAUGAG X CGAA AGAAGGGU	119	ACCCTTCT T CCGTTGAT	647
800	UAUCAACG CUGAUGAG X CGAA AAGAAGGG	120	CCCTTCTT C CGTTGATA	648
804	UUGAUUUC CUGAUGAG X CGAA ACGGAAGA	121	TCTTCCGT T GATATCAA	649
808	UUUCUUGA CUGAUGAG X CGAA AUCAACGG	122	CCGTTGAT A TCAAGAAA	650
810	ACUUUCUU CUGAUGAG X CGAA AUUAUAC	123	GTTGATAT C AAGAAAGT	651
824	UCAUUUCU CUGAUGAG X CGAA ACAGCACU	124	AGTGCTGT T AGAAATGA	652
825	CUCAUUUC CUGAUGAG X CGAA AACAGCAC	125	GTGCTGTT A GAAATGAG	653
839	CCAUCCGA CUGAUGAG X CGAA ACUCCUC	126	GAGGAAGT T TCGATGG	654
840	CCCAUCCG CUGAUGAG X CGAA AACUCCU	127	AGGAAGTT T CGATGGG	655
841	CCCAUCC CUGAUGAG X CGAA AAACUCC	128	GGAAGTTT C GGATGGG	656
855	GCUGUCUG CUGAUGAG X CGAA AUCAGCCC	129	GGGCTGAT C CAGACAGC	657
878	GGUAGGAG CUGAUGAG X CGAA AGCGCAGC	130	GCTGCGCT T CTCCTACC	658
879	AGGUAGGA CUGAUGAG X CGAA AAGCGCAG	131	CTGCGCTT C TCCTACCT	659
881	CCAGGUAG CUGAUGAG X CGAA AGAAGCGC	132	GCGCTTCT C CTACCTGG	660
884	CAGCCAGG CUGAUGAG X CGAA AGGAGAAG	133	CTTCTCCT A CCTGGCTG	661
897	GCACCUUC CUGAUGAG X CGAA AUCACAGC	134	GCTGTGAT C GAAGGTGC	662
911	CCAUGAUG CUGAUGAG X CGAA AUUUGGCA	135	TGCCAAAT T CATCATGG	663
912	CCCAUGAU CUGAUGAG X CGAA AAUUGGC	136	GCCAAATT C ATCATGGG	664
915	UCCCCAU CUGAUGAG X CGAA AUGAAUUU	137	AAATTCAT C ATGGGGGA	665
926	GCACGGAA CUGAUGAG X CGAA AGUCCCCC	138	GGGGGACT C TTCCGTGC	666

Table 3

928	CUGCACGG CUGAUGAG X CGAA AGAGUCCC	139	GGGACTCT T CCGTGCAG	667
929	CCUGCACG CUGAUGAG X CGAA AAGAGUCC	140	GGACTCTT C CGTGCAGG	668
940	CUUCCACU CUGAUGAG X CGAA AUCCUGCA	141	TGCAGGAT C AGTGGAAAG	669
954	UCGUGGGA CUGAUGAG X CGAA AGCUCCUU	142	AAGGAGCT T TCCCACGA	670
955	CUCGUGGG CUGAUGAG X CGAA AAGCUCCU	143	AGGAGCTT T CCCACGAG	671
956	CCUCGUGG CUGAUGAG X CGAA AAAGCUCC	144	GGAGCTTT C CCACGAGG	672
988	UGGGGGGA CUGAUGAG X CGAA AUGCUCGG	145	CCGAGCAT A TCCCCCA	673
990	GGUGGGGG CUGAUGAG X CGAA AUAUGCUC	146	GAGCATAT C CCCCCACC	674
1000	UGGCCGGG CUGAUGAG X CGAA AGGUGGGG	147	CCCCACCT C CCCGGCCA	675
1020	GGCUCCAG CUGAUGAG X CGAA AUUCGUUU	148	AAACGAAT C CTGGAGCC	676
1052	UUGGGAAG CUGAUGAG X CGAA ACUCCCUG	149	CAGGGAGT T CTTCCTAA	677
1053	UUUGGGAA CUGAUGAG X CGAA AACUCCCU	150	AGGGAGTT C TTCCCAA	678
1055	GAUUUGGG CUGAUGAG X CGAA AGAACUCC	151	GGAGTTCT T CCCAAATC	679
1056	UGAUUUGG CUGAUGAG X CGAA AAGAACUC	152	GAGTTCTT C CCAAATCA	680
1063	CCACUGGU CUGAUGAG X CGAA AUUUGGGA	153	TCCCAAAT C ACCAGTGG	681
1096	GCAGUCUU CUGAUGAG X CGAA AUCCUCCU	154	AGGAGGAT A AAGACTGC	682
1110	UCUUCUUU CUGAUGAG X CGAA AUGGGGCA	155	TGCCCCAT C AAGGAAGA	683
1133	CGGCAUUU CUGAUGAG X CGAA AGGGGCUU	156	AAGCCCCT T AAATGCCG	684
1134	GCGGCAUU CUGAUGAG X CGAA AAGGGGCU	157	AGCCCCTT A AATGCCGC	685
1148	CGAUGCCG CUGAUGAG X CGAA AGGGUGCG	158	CGCACCCCT A CGGCATCG	686
1155	AUGCUUUC CUGAUGAG X CGAA AUGCCGUA	159	TACGGCAT C GAAAGCAT	687
1168	AGUGUCUU CUGAUGAG X CGAA ACUCAUGC	160	GCATGAGT C AAGACACT	688
1182	CGACUUCU CUGAUGAG X CGAA ACUUCAGU	161	ACTGAAGT T AGAAGTCG	689
1183	CCGACUUC CUGAUGAG X CGAA AACUUCAG	162	CTGAAGTT A GAAGTCGG	690
1189	CACGACCC CUGAUGAG X CGAA ACUUCUAA	163	TTAGAAGT C GGGTCGTG	691
1194	CCCCCCAC CUGAUGAG X CGAA ACCCGACU	164	AGTCGGGT C GTGGGGGG	692
1207	ACCUCGAA CUGAUGAG X CGAA ACUUCCCC	165	GGGGAAGT C TTCGAGGT	693
1209	GCACCUCG CUGAUGAG X CGAA AGACUUC	166	GGAGTCTT T CGAGGTGC	694
1210	GGCACCUC CUGAUGAG X CGAA AAGACUUC	167	GAAGTCTT C GAGGTGCC	695
1229	UGGUGGGG CUGAUGAG X CGAA AGGCAGCC	168	GGCTGCCT C CCCAGCCA	696
1250	CGGGCAGU CUGAUGAG X CGAA ACGGCUCC	169	GGAGCCGT C ACTGCCCG	697
1285	CUUCCAGU CUGAUGAG X CGAA ACUCAGUG	170	CACTGAGT T ACTGGAAG	698
1286	GCUUCCAG CUGAUGAG X CGAA AACUCAGU	171	ACTGAGTT A CTGGAAGC	699
1298	UGACCAGG CUGAUGAG X CGAA AGGGCUUC	172	GAAGCCCT T CCTGGTCA	700
1299	UUGACCAG CUGAUGAG X CGAA AAGGGCUU	173	AAGCCCTT C CTGGTCAA	701
1305	CACAUGUU CUGAUGAG X CGAA ACCAGGAA	174	TTCTTGGT C AACATGTG	702
1321	GAGGACCG CUGAUGAG X CGAA AGCCACGC	175	GCGTGGCT A CGGTCCTC	703
1326	GCCGUGAG CUGAUGAG X CGAA ACCGUAGC	176	GCTACGGT C CTCACGGC	704
1329	CCGGCCGU CUGAUGAG X CGAA AGGACCGU	177	ACGGTCCT C ACGGCCGG	705
1342	GCAGAGGU CUGAUGAG X CGAA AGCGCCGG	178	CCGGCGCT T ACCTCTGC	706
1343	AGCAGAGG CUGAUGAG X CGAA AAGCGCCG	179	CGGCGCTT A CCTCTGCT	707
1347	CUGUAGCA CUGAUGAG X CGAA AGGUAAGC	180	GCTTACCT C TGCTACAG	708
1352	GGAACCUG CUGAUGAG X CGAA AGCAGAGG	181	CCTCTGCT A CAGGTTCC	709
1358	UGAACAGG CUGAUGAG X CGAA ACCUGUAG	182	CTACAGGT T CCTGTTCA	710
1359	UUGAACAG CUGAUGAG X CGAA AACCUGUA	183	TACAGGTT C CTGTTCAA	711
1364	UGCUGUUG CUGAUGAG X CGAA ACAGGAAC	184	GTTCTGTG T CAACAGCA	712
1365	UUGCUGUU CUGAUGAG X CGAA AACAGGAA	185	TTCTGTGT C AACAGCAA	713

Table 3

1379	GGUCAGGC CUGAUGAG X CGAA AUGUGUUG	186	CAACACAT A GCCTGACC	714
1390	GAGUGGAG CUGAUGAG X CGAA AGGGUCAG	187	CTGACCCT C CTCCACTC	715
1393	GUGGAGUG CUGAUGAG X CGAA AGGAGGGU	188	ACCCTCCT C CACTCCAC	716
1398	UGGAGGUG CUGAUGAG X CGAA AGUGGAGG	189	CCTCCACT C CACCTCCA	717
1404	AGUGGGUG CUGAUGAG X CGAA AGGUGGAG	190	CTCCACCT C CACCCACT	718
1415	CAGAGGCG CUGAUGAG X CGAA ACAGUGGG	191	CCCCTGT C CGCTCTG	719
1421	UGCGGGCA CUGAUGAG X CGAA AGGCGGAC	192	GTCCGCCT C TGCCGCA	720
1446	AUGCCUGC CUGAUGAG X CGAA AGUCGGGC	193	GCCCGACT A GCAGGCAT	721
1463	CCCUUACC CUGAUGAG X CGAA ACCGCGGC	194	GCCGCGT A GGTAAGGG	722
1467	GCGGCCCU CUGAUGAG X CGAA ACCUACCG	195	CGGTAGGT A AGGGCCGC	723
1486	CGGCUCUC CUGAUGAG X CGAA ACGCGGUC	196	GACCGCGT A GAGAGCG	724
1511	GCAGAACC CUGAUGAG X CGAA ACGUCCGU	197	ACGGACGT T GGTCTGC	725
1515	UAGUGCAG CUGAUGAG X CGAA ACCAACGU	198	ACGTTGGT T CTGACTA	726
1516	UUAGUGCA CUGAUGAG X CGAA AACCAACG	199	CGTTGGTT C TGCACTAA	727
1523	AUGGGUUU CUGAUGAG X CGAA AGUGCAGA	200	TCTGCACT A AAACCCAT	728
1532	CCGGGGAA CUGAUGAG X CGAA AUGGGUUU	201	AAACCCAT C TTCCCCGG	729
1534	AUCCGGGG CUGAUGAG X CGAA AGAUGGGU	202	ACCCATCT T CCCCAGAT	730
1535	CAUCCGGG CUGAUGAG X CGAA AAGAUGGG	203	CCCCTCTT C CCCGATG	731
1549	AGGGGUGA CUGAUGAG X CGAA ACACACAU	204	ATGTGTGT C TCACCCCT	732
1551	UGAGGGGU CUGAUGAG X CGAA AGACACAC	205	GTGTGTCT C ACCCCTCA	733
1558	AAAAGGAU CUGAUGAG X CGAA AGGGGUGA	206	TCACCCCT C ATCTTTT	734
1561	AGUAAAAG CUGAUGAG X CGAA AUGAGGGG	207	CCCCTCAT C CTTTACT	735
1564	AAAAGUAA CUGAUGAG X CGAA AGGAUGAG	208	CTCATCCT T TTACTTTT	736
1565	AAAAAGUA CUGAUGAG X CGAA AAGGAUGA	209	TCATCCTT T TACTTTT	737
1566	CAAAAAGU CUGAUGAG X CGAA AAAGGAUG	210	CATCCTTT T ACTTTTG	738
1567	GCAAAAAG CUGAUGAG X CGAA AAAAGGAU	211	ATCCTTTT A CTTTTTG	739
1570	GGGGCAAA CUGAUGAG X CGAA AGUAAAAG	212	CTTTACT T TTTGCCCC	740
1571	AGGGGCAA CUGAUGAG X CGAA AAGUAAAA	213	TTTTACTT T TTGCCCT	741
1572	AAGGGGCA CUGAUGAG X CGAA AAAGUAAA	214	TTACTTTT T TGCCCTT	742
1573	GAAGGGGC CUGAUGAG X CGAA AAAAGUAA	215	TTACTTTT T GCCCTTC	743
1580	CAAAGUGG CUGAUGAG X CGAA AGGGGCAA	216	TGCCCCCT T CCACTTG	744
1581	UCAAGUG CUGAUGAG X CGAA AAGGGGCA	217	TGCCCCCT C CACTTTGA	745
1586	GGUACUCA CUGAUGAG X CGAA AGUGGAAG	218	CTTCCACT T TGAGTACC	746
1587	UGGUACUC CUGAUGAG X CGAA AAGUGGAA	219	TTCCACTT T GAGTACCA	747
1592	GGAUUUGG CUGAUGAG X CGAA ACUCAAG	220	CTTTGAGT A CCAAATCC	748
1599	GGCUUGUG CUGAUGAG X CGAA AUUUGGUA	221	TACCAAAT C CACAAGCC	749
1610	CCUCAAAA CUGAUGAG X CGAA AUGGCUUG	222	CAAGCCAT T TTTGAGG	750
1611	UCCUCAAA CUGAUGAG X CGAA AAUGGCUU	223	AAGCCATT T TTTGAGGA	751
1612	CUCUCAAA CUGAUGAG X CGAA AAAUGGCU	224	AGCCATTT T TTGAGGAG	752
1613	UCUCCUCA CUGAUGAG X CGAA AAAAUGGC	225	GCCATTTT T TGAGGAGA	753
1614	CUCUCCUC CUGAUGAG X CGAA AAAAUGG	226	CCATTTTT T GAGGAGAG	754
1634	CAGCAUGG CUGAUGAG X CGAA ACUCUCUU	227	AAGAGAGT A CCATGCTG	755
1665	GACGGGUG CUGAUGAG X CGAA AGGCCCUU	228	AGGGGCCT A CACCCGTC	756
1673	AGCCCCAA CUGAUGAG X CGAA ACGGGUGU	229	ACACCCGT C TTGGGGCT	757
1675	CGAGCCCC CUGAUGAG X CGAA AGACGGGU	230	ACCCGTCT T GGGGCTCG	758
1682	GGUGGGGC CUGAUGAG X CGAA AGCCCCAA	231	TTGGGGCT C GCCCACC	759
1698	CCAGGAGG CUGAUGAG X CGAA AGCCUGG	232	CCAGGGCT C CCTCCTGG	760

Table 3

1702	UGCUCAG CUGAUGAG X CGAA AGGGAGCC	233	GGCTCCCT C CTGGAGCA	761
1712	CCGCCUGG CUGAUGAG X CGAA AUGCUCCA	234	TGGAGCAT C CCAGGCGG	762
1746	GCAGAUUC CUGAUGAG X CGAA AGGGGGGG	235	CCCCCCT T GAATCTGC	763
1751	UCCUGCA CUGAUGAG X CGAA AUUCAAGG	236	CCTTGAAT C TGCAGGGA	764
1766	GGAGUGGA CUGAUGAG X CGAA AGUUGCUC	237	GAGCAACT C TCCACTCC	765
1768	AUGGAGUG CUGAUGAG X CGAA AGAGUUGC	238	GCAACTCT C CACTCCAT	766
1773	UAAUAUG CUGAUGAG X CGAA AGUGGAGA	239	TCTCCACT C CATATTTA	767
1777	UAAUAAA CUGAUGAG X CGAA AUGGAGUG	240	CACTCCAT A TTTATTTA	768
1779	UUUAAUA CUGAUGAG X CGAA AUUUGGAG	241	CTCCATAT T TATTTAAA	769
1780	GUUUAUU CUGAUGAG X CGAA AAUAUGGA	242	TCCATATT T ATTTAAAC	770
1781	UGUUUAAA CUGAUGAG X CGAA AAUAUGG	243	CCATATT A TTTAAACA	771
1783	AUUGUUUA CUGAUGAG X CGAA AUAAAUAU	244	ATATTTAT T TAAACAAT	772
1784	AAUUGUUU CUGAUGAG X CGAA AAUAAAUA	245	TATTTAT T AAACAATT	773
1785	AAAUUGUU CUGAUGAG X CGAA AAUAAAUA	246	ATTTATTT A AACAATTT	774
1792	GGGGAAA CUGAUGAG X CGAA AUUGUUUA	247	TAAACAAT T TTTCCCC	775
1793	UGGGGAAA CUGAUGAG X CGAA AAUUGUUU	248	AAACAATT T TTTCCCCA	776
1794	UUGGGGAA CUGAUGAG X CGAA AAUUGUUU	249	AACAATTT T TTCCCCAA	777
1795	UUUGGGGA CUGAUGAG X CGAA AAAAUUGU	250	ACAATTTT T TCCCCAAA	778
1796	CUUUGGGG CUGAUGAG X CGAA AAAAUUG	251	CAATTTT T CCCCCAAG	779
1797	CCUUGGG CUGAUGAG X CGAA AAAAUUU	252	AATTTT C CCCCCAAGG	780
1809	GCACUAUG CUGAUGAG X CGAA AUGCCUUU	253	AAAGGCAT C CATAGTGC	781
1813	UAGUGCAC CUGAUGAG X CGAA AUGGAUGC	254	GCATCCAT A GTGCACTA	782
1821	GAAAUGC CUGAUGAG X CGAA AGUGCACU	255	AGTGCACT A GCATTTTC	783
1826	UUCAAGAA CUGAUGAG X CGAA AUGCUAGU	256	ACTAGCAT T TTCTTGAA	784
1827	GUUCAAGA CUGAUGAG X CGAA AAUGCUG	257	CTAGCATT T TCTGAAC	785
1828	GGUUCAAG CUGAUGAG X CGAA AAUGCUGA	258	TAGCATT T CTTGAACC	786
1829	UGGUUCA CUGAUGAG X CGAA AAAAUGC	259	AGCATTT C TTGAACCA	787
1831	AUUGGUUC CUGAUGAG X CGAA AGAAAUG	260	CATTTCT T GAACCAAT	788
1840	UAAUACAU CUGAUGAG X CGAA AUUGGUUC	261	GAACCAAT A ATGTATTA	789
1845	AAUUUUA CUGAUGAG X CGAA ACAUUUAU	262	AATAATGT A TTAATAAT	790
1847	AAAAUUU CUGAUGAG X CGAA AUACAUUA	263	TAATGTAT T AAAATTTT	791
1848	AAAAUUU CUGAUGAG X CGAA AAUACAUU	264	AATGTAT A AAATTTT	792
1853	CAUCAAAA CUGAUGAG X CGAA AUUUUAU	265	ATTAAAT T TTTGATG	793
1854	ACAUCAAA CUGAUGAG X CGAA AAUUUUA	266	TTAAAT T TTTGATGT	794
1855	GACAUCAA CUGAUGAG X CGAA AAUUUUA	267	TAAATTT T TTGATGTC	795
1856	UGACAUCA CUGAUGAG X CGAA AAAAUUU	268	AAAATTTT T TGATGTCA	796
1857	CUGACAUC CUGAUGAG X CGAA AAAAUUU	269	AAATTTT T GATGTGAG	797
1863	GCAAGGCU CUGAUGAG X CGAA ACAUCAAA	270	TTGATGT C AGCCTTGC	798
1869	CUUGAUGC CUGAUGAG X CGAA AGGCUGAC	271	GTCAGCCT T GCATCAAG	799
1874	AAGCCCUU CUGAUGAG X CGAA AUGCAAGG	272	CCTTGCAT C AAGGGCTT	800
1882	UUUUGAUA CUGAUGAG X CGAA AGCCCUUG	273	CAAGGGCT T TATCAAAA	801
1883	UUUUGAU CUGAUGAG X CGAA AAGCCCUU	274	AAGGGCTT T ATCAAAAA	802
1884	CUUUUGA CUGAUGAG X CGAA AAAGCCCU	275	AGGGCTT A TCAAAAAG	803
1886	UACUUUUU CUGAUGAG X CGAA AUAAAGCC	276	GGCTTTAT C AAAAAGTA	804
1894	UAUUUAUG CUGAUGAG X CGAA ACUUUUUG	277	CAAAAAGT A CAATAATA	805
1899	GGAUUUAU CUGAUGAG X CGAA AUUGUACU	278	AGTACAAT A ATAAATCC	806
1902	UGAGGAUU CUGAUGAG X CGAA AUUAUUGU	279	ACAATAAT A AATCCTCA	807

Table 3

1906	UACCUGAG CUGAUGAG X CGAA AUUUAUUA	280	TAATAAAT C CTCAGGTA	808
1909	UACUACCU CUGAUGAG X CGAA AGGAUUUA	281	TAAATCCT C AGGTAGTA	809
1914	CCCAGUAC CUGAUGAG X CGAA ACCUGAGG	282	CCTCAGGT A GTACTGGG	810
1917	AUUCCCAG CUGAUGAG X CGAA ACUACCUG	283	CAGGTAGT A CTGGGAAT	811
1934	CCAUGGCA CUGAUGAG X CGAA AGCCUUC	284	GGAAGGCT T TGCCATGG	812
1935	CCCAUGGC CUGAUGAG X CGAA AAGCCUUC	285	GAAGGCTT T GCCATGGG	813
1954	ACUGGUCU CUGAUGAG X CGAA ACGCAGCA	286	TGCTGCGT C AGACCAGT	814
1963	CUUCCAG CUGAUGAG X CGAA ACUGGUCU	287	AGACCAGT A CTGGGAAG	815
1981	CUGCUUAC CUGAUGAG X CGAA ACCGUCCU	288	AGGACGGT T GTAAGCAG	816
1984	CAACUGCU CUGAUGAG X CGAA ACAACCGU	289	ACGTTGT A AGCAGTTG	817
1991	UAAAUAA CUGAUGAG X CGAA ACUGCUUA	290	TAAGCAGT T GTTATTTA	818
1994	CACUAAAU CUGAUGAG X CGAA ACAACUGC	291	GCAGTTGT T ATTTAGTG	819
1995	UCACUAAA CUGAUGAG X CGAA AACACUG	292	CAGTTGTT A TTTAGTGA	820
1997	UAUCACUA CUGAUGAG X CGAA AUAACAAC	293	GTGTTAT T TAGTGATA	821
1998	AUAUCACU CUGAUGAG X CGAA AAUAACAA	294	TTGTTATT T AGTGATAT	822
1999	AAUAUCAC CUGAUGAG X CGAA AAUAACA	295	TGTTATTT A GTGATATT	823
2005	ACCCACAA CUGAUGAG X CGAA AUCACUAA	296	TTAGTGAT A TTGTGGGT	824
2007	UUACCCAC CUGAUGAG X CGAA AUAUCACU	297	AGTGATAT T GTGGGTAA	825
2014	UCUCACGU CUGAUGAG X CGAA ACCACAA	298	TTGTGGGT A ACGTGAGA	826
2027	CAUUGUUC CUGAUGAG X CGAA AUCUUCUC	299	GAGAAGAT A GAACAATG	827
2038	AUAUAUUA CUGAUGAG X CGAA AGCAUUGU	300	ACAATGCT A TAATATAT	828
2040	UUAUAUUA CUGAUGAG X CGAA AUAGCAUU	301	AATGCTAT A ATATATAA	829
2043	UCAUUAUA CUGAUGAG X CGAA AUUAUAGC	302	GCTATAAT A TATAATGA	830
2045	GUUCAUUA CUGAUGAG X CGAA AUUAUUA	303	TATAATAT A TAATGAAC	831
2047	GUGUUCAU CUGAUGAG X CGAA AUAUAUUA	304	TAATATAT A ATGAACAC	832
2062	UUAUUAAA CUGAUGAG X CGAA ACCCACGU	305	ACGTGGGT A TTTAATAA	833
2064	UCUUAUUA CUGAUGAG X CGAA AUACCCAC	306	GTGGGTAT T TAATAAGA	834
2065	UUCUUAUU CUGAUGAG X CGAA AAUACCCA	307	TGGGTATT T AATAAGAA	835
2066	UUUCUUAU CUGAUGAG X CGAA AAAUACCC	308	GGGTATTT A ATAAGAAA	836
2069	AUGUUUCU CUGAUGAG X CGAA AUUAAUUA	309	TATTTAAT A AGAAACAT	837
2088	GACAAAGU CUGAUGAG X CGAA AUCUCACA	310	TGTGAGAT T ACTTTGTC	838
2089	GGACAAAG CUGAUGAG X CGAA AAUCUCAC	311	GTGAGATT A CTTTGTC	839
2092	GCGGGACA CUGAUGAG X CGAA AGUAAUCU	312	AGATTACT T TGTCCCGC	840
2093	AGCGGGAC CUGAUGAG X CGAA AAGUAAUC	313	GATTACTT T GTCCCGCT	841
2096	AUAAGCGG CUGAUGAG X CGAA ACAAAGUA	314	TACTTTGT C CCGCTTAT	842
2102	AGCAGAAU CUGAUGAG X CGAA AGCGGGAC	315	GTCCCGCT T ATTCTGCT	843
2103	GAGCAGAA CUGAUGAG X CGAA AAGCGGGA	316	TCCCGCTT A TTCTGCTC	844
2105	GGGAGCAG CUGAUGAG X CGAA AUAAGCGG	317	CCGCTTAT T CTGCTCCC	845
2106	AGGGAGCA CUGAUGAG X CGAA AAUAAGCG	318	CGCTTATT C TGCTCCCT	846
2111	AUAACAGG CUGAUGAG X CGAA AGCAGAAU	319	ATTCTGCT C CCTGTTAT	847
2117	UAGCAGAU CUGAUGAG X CGAA ACAGGGAG	320	CTCCCTGT T ATCTGCTA	848
2118	CUAGCAGA CUGAUGAG X CGAA AACAGGGA	321	TCCCTGTT A TCTGCTAG	849
2120	AUCUAGCA CUGAUGAG X CGAA AUAACAGG	322	CCTGTTAT C TGCTAGAT	850
2125	ACUAGAUC CUGAUGAG X CGAA AGCAGAU	323	TATCTGCT A GATCTAGT	851
2129	GAGAACUA CUGAUGAG X CGAA AUCUAGCA	324	TGCTAGAT C TAGTTCTC	852
2131	UUGAGAAC CUGAUGAG X CGAA AGAUCUAG	325	CTAGATCT A GTTCTCAA	853
2134	UGAUUGAG CUGAUGAG X CGAA ACUAGAUC	326	GATCTAGT T CTCAATCA	854

Table 3

2135	GUGAUUGA CUGAUGAG X CGAA AACUAGAU	327	ATCTAGTT C TCAATCAC	855
2137	CAGUGAUU CUGAUGAG X CGAA AGAACUAG	328	CTAGTTCT C AATCACTG	856
2141	GGAGCAGU CUGAUGAG X CGAA AUUGAGAA	329	TTCTCAAT C ACTGCTCC	857
2148	ACACGGGG CUGAUGAG X CGAA AGCAGUGA	330	TCACTGCT C CCCCGTGT	858
2159	CAUUCUAA CUGAUGAG X CGAA ACACACGG	331	CCGTGTGT A TTAGAATG	859
2161	UGCAUUCU CUGAUGAG X CGAA AUACACAC	332	GTGTGTAT T AGAATGCA	860
2162	AUGCAUUC CUGAUGAG X CGAA AAUACACA	333	TGTGTATT A GAATGCAT	861
2173	GAAGACCU CUGAUGAG X CGAA ACAUGCAU	334	ATGCATGT A AGGTCTTC	862
2178	CACAAGAA CUGAUGAG X CGAA ACCUUAUA	335	TGTAAGGT C TTCTGTGT	863
2180	GACACAAG CUGAUGAG X CGAA AGACCUUA	336	TAAGGTCT T CTGTGTCT	864
2181	GGACACAA CUGAUGAG X CGAA AAGACCUU	337	AAGGTCTT C TTGTGTCC	865
2183	CAGGACAC CUGAUGAG X CGAA AGAAGACC	338	GGTCTTCT T GTGTCTTG	866
2188	UUCAUCAG CUGAUGAG X CGAA ACACAAGA	339	TCTGTGT C CTGATGAA	867
2201	CAAGCACA CUGAUGAG X CGAA AUUUUUA	340	TGAAAAAT A TGTGCTTG	868
2208	CUCAUUUC CUGAUGAG X CGAA AGCACAUUA	341	TATGTGCT T GAAATGAG	869
2222	AGAGAUCA CUGAUGAG X CGAA AGUUUCUC	342	GAGAAACT T TGATCTCT	870
2223	CAGAGAU CUGAUGAG X CGAA AAGUUUCU	343	AGAAACTT T GATCTCTG	871
2227	UAAGCAGA CUGAUGAG X CGAA AUCAAAGU	344	ACTTGAT C TCTGCTTA	872
2229	AGUAAGCA CUGAUGAG X CGAA AGAUCAAA	345	TTGATCT C TGCTTACT	873
2234	ACAUIUAGU CUGAUGAG X CGAA AGCAGAGA	346	TCTCTGCT T ACTAATGT	874
2235	CACAUUAG CUGAUGAG X CGAA AAGCAGAG	347	CTCTGCTT A CTAATGTG	875
2238	GGGCACAU CUGAUGAG X CGAA AGUAAGCA	348	TGCTTACT A ATGTGCCC	876
2252	UGGACUUG CUGAUGAG X CGAA ACAUGGGG	349	CCCATGT C CAAGTCCA	877
2258	GCAGGUUG CUGAUGAG X CGAA ACUUGGAC	350	GTCCAAGT C CAACCTGC	878
2283	CAUGUAAU CUGAUGAG X CGAA AUCAGGUC	351	GACCTGAT C ATTACATG	879
2286	AGCCAUGU CUGAUGAG X CGAA AUGAUCAG	352	CTGATCAT T ACATGGCT	880
2287	CAGCCAUG CUGAUGAG X CGAA AAUGAUCA	353	TGATCATT A CATGGCTG	881
2300	GGCUUAGG CUGAUGAG X CGAA ACCACAGC	354	GCTGTGGT T CCTAAGCC	882
2301	AGGCUUAG CUGAUGAG X CGAA AACCACAG	355	CTGTGGTT C CTAAGCCT	883
2304	AACAGGCU CUGAUGAG X CGAA AGGAACCA	356	TGGTTCCT A AGCCTGTT	884
2312	ACUUCAGC CUGAUGAG X CGAA ACAGGCUU	357	AAGCCTGT T GCTGAAGT	885
2321	GCGACAAU CUGAUGAG X CGAA ACUUCAGC	358	GCTGAAGT C ATTGTGCG	886
2324	UGAGCGAC CUGAUGAG X CGAA AUGACUUC	359	GAAGTCAT T GTCGCTCA	887
2327	UGCUGAGC CUGAUGAG X CGAA ACAAUGAC	360	GTCATTGT C GCTCAGCA	888
2331	CUAUUGCU CUGAUGAG X CGAA AGCGACAA	361	TTGTGCT C AGCAATAG	889
2338	CUGCACCC CUGAUGAG X CGAA AUUGCUGA	362	TCAGCAAT A GGGTGCAG	890
2348	UCCUGGAA CUGAUGAG X CGAA ACUGCACC	363	GGTGCAGT T TTCCAGGA	891
2349	UUCCUGGA CUGAUGAG X CGAA AACUGCAC	364	GTGCAGTT T TCCAGGAA	892
2350	AUUCUGG CUGAUGAG X CGAA AAACUGCA	365	TGCAGTTT T CCAGGAAT	893
2351	UAUUCUG CUGAUGAG X CGAA AAAACUGC	366	GCAGTTT C CAGGAATA	894
2359	CAAAUGCC CUGAUGAG X CGAA AUUCUGG	367	CCAGGAAT A GGCATTGT	895
2365	AUUAGGCA CUGAUGAG X CGAA AUGCCUAU	368	ATAGGCAT T TGCCTAAT	896
2366	AAUUAGGC CUGAUGAG X CGAA AAUGCCUA	369	TAGGCATT T GCCTAATT	897
2371	CCAGGAAU CUGAUGAG X CGAA AGGCAAU	370	ATTGCTT A ATTCTGCT	898
2374	AUGCCAGG CUGAUGAG X CGAA AUUAGGCA	371	TGCCTAAT T CCTGGCAT	899
2375	CAUGCCAG CUGAUGAG X CGAA AAUAGGC	372	GCCTAATT C CTGGCATG	900
2389	AGUCACUA CUGAUGAG X CGAA AGUGUCAU	373	ATGACACT C TAGTGACT	901

Table 3

2391	GAAGUCAC CUGAUGAG X CGAA AGAGUGUC	374	GACACTCT A GTGACTTC	902
2398	UCACCAGG CUGAUGAG X CGAA AGUCACUA	375	TAGTGACT T CTTGGTGA	903
2399	CUCACCAG CUGAUGAG X CGAA AAGUCACU	376	AGTGACTT C CTGGTGAG	904
2419	UGUACCAG CUGAUGAG X CGAA ACAGGCUG	377	CAGCCTGT C CTGGTACA	905
2425	CCCUGCUG CUGAUGAG X CGAA ACCAGGAC	378	GTCCTGGT A CAGCAGGG	906
2435	UACAGCAA CUGAUGAG X CGAA ACCCUGCU	379	AGCAGGGT C TTGCTGTA	907
2437	GUUACAGC CUGAUGAG X CGAA AGACCCUG	380	CAGGGTCT T GCTGTAAC	908
2443	GUCUGAGU CUGAUGAG X CGAA ACAGCAAG	381	CTTGCTGT A ACTCAGAC	909
2447	GAAUGUCU CUGAUGAG X CGAA AGUUCACAG	382	CTGTAAC T C AGACATTC	910
2454	ACCCUUGG CUGAUGAG X CGAA AUGUCUGA	383	TCAGACAT T CCAAGGGT	911
2455	UACCCUUG CUGAUGAG X CGAA AAUGUCUG	384	CAGACATT C CAAGGGTA	912
2463	GCUCCCCA CUGAUGAG X CGAA ACCCUUGG	385	CCAAGGGT A TGGGAAGC	913
2475	GGUGUGAA CUGAUGAG X CGAA AUGGCUUC	386	GAAGCCAT A TTCACACC	914
2477	GAGGUGUG CUGAUGAG X CGAA AUAUGGCU	387	AGCCATAT T CACACCTC	915
2478	UGAGGUGU CUGAUGAG X CGAA AAUAUGGC	388	GCCATATT C ACACCTCA	916
2485	CAGAGCGU CUGAUGAG X CGAA AGGUGUGA	389	TCACACCT C ACGCTCTG	917
2491	CAUGUCCA CUGAUGAG X CGAA AGCGUGAG	390	CTCACGCT C TGGACATG	918
2502	CUUCCCUA CUGAUGAG X CGAA AUCAUGUC	391	GACATGAT T TAGGGAAG	919
2503	GCUCCCCU CUGAUGAG X CGAA AAUCAUGU	392	ACATGATT T AGGGAAGC	920
2504	UGCUCUCC CUGAUGAG X CGAA AAAUCAUG	393	CATGATTT A GGGGAAGCA	921
2536	UGAUCCCA CUGAUGAG X CGAA AGGUGGGG	394	CCCCACCT T TGGGATCA	922
2537	CUGAUCCC CUGAUGAG X CGAA AAGUGGGG	395	CCCACCTT T GGGATCAG	923
2543	CGGAGGCU CUGAUGAG X CGAA AUCCCAAA	396	TTTGGGAT C AGCCTCCG	924
2549	GAAUGGCG CUGAUGAG X CGAA AGGCUGAU	397	ATCAGCCT C CGCCATTC	925
2556	CGACUUGG CUGAUGAG X CGAA AUGGCGGA	398	TCCGCCAT T CCAAGTCG	926
2557	UCGACUUG CUGAUGAG X CGAA AAUGGCGG	399	CCGCCATT C CAAGTCGA	927
2563	AGAGUGUC CUGAUGAG X CGAA ACUUGGAA	400	TTCCAAGT C GACACTCT	928
2570	CUCAAGAA CUGAUGAG X CGAA AGUGUCGA	401	TCGACACT C TTCTTGAG	929
2572	UGCUC AAG CUGAUGAG X CGAA AGAGUGUC	402	GACACTCT T CTTGAGCA	930
2573	CUGCUC AA CUGAUGAG X CGAA AAGAGUGU	403	ACACTCTT C TTGAGCAG	931
2575	GUCUGCUC CUGAUGAG X CGAA AGAAGAGU	404	ACTCTTCT T GAGCAGAC	932
2590	CUCUCCA CUGAUGAG X CGAA AUCACGGU	405	ACCGTGAT T TGGGAAGAG	933
2591	UCUCUCC CUGAUGAG X CGAA AAUCACGG	406	CCGTGATT T GGAAGAGA	934
2622	GUUUC AAG CUGAUGAG X CGAA AGUGUGGU	407	ACCACACT T CTTGAAAC	935
2623	UGUUUCAA CUGAUGAG X CGAA AAGUGUGG	408	CCACACTT C TTGAAACA	936
2625	GCUGUUUC CUGAUGAG X CGAA AGAAGUGU	409	ACACTTCT T GAAACAGC	937
2646	GCCUAAAG CUGAUGAG X CGAA ACCGUCAC	410	GTGACGGT C CTTTAGGC	938
2649	GCUGCCUA CUGAUGAG X CGAA AGGACCGU	411	ACGGTCCT T TAGGCAGC	939
2650	GGCUGCCU CUGAUGAG X CGAA AAGGACCG	412	CGGTCCTT T AGGCAGCC	940
2651	AGGUGGCC CUGAUGAG X CGAA AAAGGACC	413	GGTCCTTT A GGCAGCCT	941
2668	GGGACAGA CUGAUGAG X CGAA ACGGCGGC	414	GCCGCCGT C TCTGTCCC	942
2670	CCGGGACA CUGAUGAG X CGAA AGACGGCG	415	CGCCGTCT C TGTCCCGG	943
2674	UGAACCGG CUGAUGAG X CGAA ACAGAGAC	416	GTCTCTGT C CCGGTTCA	944
2680	GCAAGGUG CUGAUGAG X CGAA ACCGGGAC	417	GTCCCGGT T CACCTTGC	945
2681	GGCAAGGU CUGAUGAG X CGAA AACC GGGA	418	TCCCGGTT C ACCTTGCC	946
2686	CUCUCGGC CUGAUGAG X CGAA AGGUGAAC	419	GTTCACTT T GCCGAGAG	947
2703	GUGGGGCA CUGAUGAG X CGAA ACGCGCCU	420	AGGCGCGT C TGCCCCAC	948

Table 3

2715	CAGGGUUU CUGAUGAG X CGAA AGGGUGGG	421	CCCACCCT C AAACCCTG	949
2741	AGAGUCGU CUGAUGAG X CGAA AGCACCAU	422	ATGGTGCT C ACGACTCT	950
2748	UGCAGGAA CUGAUGAG X CGAA AGUCGUGA	423	TCACGACT C TTCCTGCA	951
2750	UUUGCAGG CUGAUGAG X CGAA AGAGUCGU	424	ACGACTCT T CCTGCAAA	952
2751	CUUUGCAG CUGAUGAG X CGAA AAGAGUCG	425	CGACTCTT C CTGCAAAG	953
2774	UUA AUGUG CUGAUGAG X CGAA AGGUCUUC	426	GAAGACCT C CACATTAA	954
2780	AGCCACUU CUGAUGAG X CGAA AUGUGGAG	427	CTCCACAT T AAGTGGCT	955
2781	AAGCCACU CUGAUGAG X CGAA AAUGUGGA	428	TCCACATT A AGTGGCTT	956
2789	AUGUUAAA CUGAUGAG X CGAA AGCCACUU	429	AAGTGGCT T TTTAACAT	957
2790	CAUGUAAA CUGAUGAG X CGAA AAGCCACU	430	AGTGGCTT T TTAACATG	958
2791	UCAUGUUA CUGAUGAG X CGAA AAAGCCAC	431	GTGGCTTT T TAACATGA	959
2792	UUCAUGUU CUGAUGAG X CGAA AAAAGCCA	432	TGGCTTTT T AACATGAA	960
2793	UUUCAUGU CUGAUGAG X CGAA AAAAAGCC	433	GGCTTTTT A ACATGAAA	961
2816	UCGGGAGC CUGAUGAG X CGAA ACAGCUGC	434	GCAGCTGT A GCTCCCGA	962
2820	UAGCUCGG CUGAUGAG X CGAA AGCUACAG	435	CTGTAGCT C CCGAGCTA	963
2828	CAAGAGAG CUGAUGAG X CGAA AGCUCGGG	436	CCCAGACT A CTCTCTTG	964
2831	UGGCAAGA CUGAUGAG X CGAA AGUAGCUC	437	GAGCTACT C TCTTGCCA	965
2833	GCUGGCAA CUGAUGAG X CGAA AGAGUAGC	438	GCTACTCT C TTGCCAGC	966
2835	AUGCUGGC CUGAUGAG X CGAA AGAGAGUA	439	TACTCTCT T GCCAGCAT	967
2844	AAUGUGAA CUGAUGAG X CGAA AUGCUGGC	440	GCCAGCAT T TTCACATT	968
2845	AAAUGUGA CUGAUGAG X CGAA AAUGCUGG	441	CCAGCATT T TCACATTT	969
2846	AAA AUGUG CUGAUGAG X CGAA AAAUGCUG	442	CAGCATTT T CACATTTT	970
2847	CAAAAUGU CUGAUGAG X CGAA AAAAUGCU	443	AGCATTTT C ACATTTTG	971
2852	AAAGGCAA CUGAUGAG X CGAA AUGUGAAA	444	TTTCACAT T TTGCCTTT	972
2853	GAAAGGCA CUGAUGAG X CGAA AAUGUGAA	445	TTCACATT T TGCCTTTC	973
2854	AGAAAGGC CUGAUGAG X CGAA AAAUGUGA	446	TCACATTT T GCCTTTCT	974
2859	CCACGAGA CUGAUGAG X CGAA AGGCAAAA	447	TTTGCCT T TCTCGTGG	975
2860	ACCACGAG CUGAUGAG X CGAA AAGGCAAA	448	TTTGCCTT T CTCGTGGT	976
2861	UACCACGA CUGAUGAG X CGAA AAAGGCAA	449	TTGCCTTT C TCGTGGTA	977
2863	UCUACCAC CUGAUGAG X CGAA AGAAAGGC	450	GCCTTTCT C GTGGTAGA	978
2869	CUGGCUUC CUGAUGAG X CGAA ACCACGAG	451	CTCGTGGT A GAAGCCAG	979
2879	UUUCUCUG CUGAUGAG X CGAA ACUGGCUU	452	AAGCCAGT A CAGAGAAA	980
2889	CACCACAG CUGAUGAG X CGAA AUUUCUCU	453	AGAGAAAT T CTGTGGTG	981
2890	CCACCACA CUGAUGAG X CGAA AAUUCUCU	454	GAGAAATT C TGTGGTGG	982
2905	ACACCUCG CUGAUGAG X CGAA AUGUUCCC	455	GGGAACAT T CGAGGTGT	983
2906	GACACCUC CUGAUGAG X CGAA AAUGUCC	456	GGAACATT C GAGGTGTC	984
2914	UGCAGGGU CUGAUGAG X CGAA ACACCUCG	457	CGAGGTGT C ACCCTGCA	985
2928	CCUCACCA CUGAUGAG X CGAA AGCUCUGC	458	GCAGAGCT A TGGTGAGG	986
2944	CUAAGCCU CUGAUGAG X CGAA AUCCACAC	459	GTGTGGAT A AGGCTTAG	987
2950	UGGCACCU CUGAUGAG X CGAA AGCCUUAU	460	ATAAGGCT T AGGTGCCA	988
2951	CUGGCACC CUGAUGAG X CGAA AAGCCUUA	461	TAAGGCTT A GGTGCCAG	989
2965	AGAAUGCU CUGAUGAG X CGAA ACAGCCUG	462	CAGGCTGT A AGCATTCT	990
2971	CAGCUCAG CUGAUGAG X CGAA AUGCUUAC	463	GTAAGCAT T CTGAGCTG	991
2972	CCAGCUCA CUGAUGAG X CGAA AAUGCUUA	464	TAAGCATT C TGAGCTGG	992
2983	AAAACAAC CUGAUGAG X CGAA AGCCAGCU	465	AGCTGGCT T GTTGTTTT	993
2986	UUAAAAAC CUGAUGAG X CGAA ACAAGCCA	466	TGGCTTGT T GTTTTAA	994
2989	GACUAAA CUGAUGAG X CGAA ACAACAAG	467	CTTGTGT T TTTAAGTC	995

Table 3

2990	GGACUUA CUGAUGAG X CGAA AACAAACA	468	TTGTTGTT T TTAAGTCC	996
2991	AGGACUUA CUGAUGAG X CGAA AAACAACA	469	TGTTGTTT T TAAGTCCT	997
2992	CAGGACUU CUGAUGAG X CGAA AAAACAAC	470	GTTGTTTT T AAGTCCTG	998
2993	ACAGGACU CUGAUGAG X CGAA AAAAACA	471	TTGTTTTT A AGTCCTGT	999
2997	AUAUACAG CUGAUGAG X CGAA ACUAAAA	472	TTTAAAGT C CTGTATAT	1000
3002	CAUACAUA CUGAUGAG X CGAA ACAGGACU	473	AGTCCTGT A TATGTATG	1001
3004	UACAUACA CUGAUGAG X CGAA AUACAGGA	474	TCCTGTAT A TGTATGTA	1002
3008	CUACUACA CUGAUGAG X CGAA ACAUAUAC	475	GTATATGT A TGTAGTAG	1003
3012	CAAACUAC CUGAUGAG X CGAA ACAUACAU	476	ATGTATGT A GTAGTTTG	1004
3015	ACCCAAAC CUGAUGAG X CGAA ACUACAUA	477	TATGTAGT A GTTGGGT	1005
3018	CACACCCA CUGAUGAG X CGAA ACUACUAC	478	GTAGTAGT T TGGGTGTG	1006
3019	ACACACCC CUGAUGAG X CGAA AACUACUA	479	TAGTAGTT T GGGTGTGT	1007
3028	ACUAUAUA CUGAUGAG X CGAA ACACACCC	480	GGGTGTGT A TATATAGT	1008
3030	CUACUAUA CUGAUGAG X CGAA AUACACAC	481	GTGTGTAT A TATAGTAG	1009
3032	UGCUCUA CUGAUGAG X CGAA AUAUACAC	482	GTGTATAT A TAGTAGCA	1010
3034	AAUGCUC CUGAUGAG X CGAA AUAUAUAC	483	GTATATAT A GTAGCATT	1011
3037	UGAAAUGC CUGAUGAG X CGAA ACUAUAUA	484	TATATAGT A GCATTTCA	1012
3042	CAUUUUGA CUGAUGAG X CGAA AUGCUACU	485	AGTAGCAT T TCAAATG	1013
3043	CCAUUUUG CUGAUGAG X CGAA AAUGCUC	486	GTAGCATT T CAAATGG	1014
3044	UCCAUUUU CUGAUGAG X CGAA AAAUGCUA	487	TAGCATTT C AAAATGGA	1015
3056	UAAACCAG CUGAUGAG X CGAA ACGUCCAU	488	ATGGACGT A CTGGTTTA	1016
3062	GGAGGUUA CUGAUGAG X CGAA ACCAGUAC	489	GTACTGGT T TAACCTCC	1017
3063	AGGAGGUU CUGAUGAG X CGAA AACCAGUA	490	TACTGGTT T AACCTCCT	1018
3064	UAGGAGGU CUGAUGAG X CGAA AAACCAGU	491	ACTGGTTT A ACCTCCTA	1019
3069	AAGGAUAG CUGAUGAG X CGAA AGGUUAAA	492	TTAACCT C CTATCCTT	1020
3072	UCCAAGGA CUGAUGAG X CGAA AGGAGGUU	493	AACCTCCT A TCCTGGA	1021
3074	UCUCCAAG CUGAUGAG X CGAA AUAGGAGG	494	CCTCCTAT C CTGGAGA	1022
3077	UGCUCUCC CUGAUGAG X CGAA AGGAUAGG	495	CCTATCCT T GGAGAGCA	1023
3093	AAGGUGGA CUGAUGAG X CGAA AGCCAGCU	496	AGCTGGCT C TCCACCTT	1024
3095	ACAAGGUG CUGAUGAG X CGAA AGAGCCAG	497	CTGGCTCT C CACCTTGT	1025
3101	UGUGUAAC CUGAUGAG X CGAA AGGUGGAG	498	CTCCACCT T GTTACACA	1026
3104	UAAUGUGU CUGAUGAG X CGAA ACAAGGUG	499	CACCTTGT T ACACATTA	1027
3105	AUAAUGUG CUGAUGAG X CGAA AACAAGGU	500	ACCTTGT A CACATTAT	1028
3111	UCUAACAU CUGAUGAG X CGAA AUGUGUAA	501	TTACACAT T ATGTTAGA	1029
3112	CUCUAACA CUGAUGAG X CGAA AAUGUGUA	502	TACACATT A TGTTAGAG	1030
3116	ACCUCUCU CUGAUGAG X CGAA ACAUAAUG	503	CATTATGT T AGAGAGGT	1031
3117	UACCUCUC CUGAUGAG X CGAA AACAUAU	504	ATTATGTT A GAGAGGTA	1032
3125	CAGCUCGC CUGAUGAG X CGAA ACCUCUCU	505	AGAGAGGT A GCAGAGCTG	1033
3136	ACAUAGCA CUGAUGAG X CGAA AGCAGCUC	506	GAGCTGCT C TGCTATGT	1034
3141	UAAGGACA CUGAUGAG X CGAA AGCAGAGC	507	GCTCTGCT A TGTCTTA	1035
3145	GGCUUAAG CUGAUGAG X CGAA ACAUAGCA	508	TGCTATGT C CTTAAGCC	1036
3148	AUUGGCUU CUGAUGAG X CGAA AGGACAU	509	TATGTCCT T AAGCCAAT	1037
3149	UAUUGGCU CUGAUGAG X CGAA AAGGACAU	510	ATGTCCTT A AGCCAATA	1038
3157	UGAGUAAA CUGAUGAG X CGAA AUUGGCUU	511	AAGCCAAT A TTTACTCA	1039
3159	GAUGAGUA CUGAUGAG X CGAA AUAUUGGC	512	GCCAATAT T TACTCATC	1040
3160	UGAUGAGU CUGAUGAG X CGAA AAUAUUGG	513	CCAATATT T ACTCATCA	1041
3161	CUGAUGAG CUGAUGAG X CGAA AAUAUUG	514	CAATATTT A CTCATCAG	1042

Table 3

3164	GACCUGAU CUGAUGAG X CGAA AGUAAAUA	515	TATTTACT C ATCAGGTC	1043
3167	AAUGACCU CUGAUGAG X CGAA AUGAGUAA	516	TTACTCAT C AGGTCATT	1044
3172	AAAAUAAU CUGAUGAG X CGAA ACCUGAUG	517	CATCAGGT C ATTATTTT	1045
3175	UAAAAAAU CUGAUGAG X CGAA AUGACCUG	518	CAGGTCAT T ATTTTTTA	1046
3176	GUAAAAAA CUGAUGAG X CGAA AAUGACCU	519	AGGTCATT A TTTTTTAC	1047
3178	UUGUAAAA CUGAUGAG X CGAA AUAAUGAC	520	GTCATTAT T TTTTACAA	1048
3179	AUUGUAAA CUGAUGAG X CGAA AAUAAUGA	521	TCATTATT T TTTACAAT	1049
3180	CAUUGUAA CUGAUGAG X CGAA AAUAAUG	522	CATTATTT T TTACAATG	1050
3181	CCAUUGUA CUGAUGAG X CGAA AAAAUAAU	523	ATTATTTT T TACAATGG	1051
3182	GCCAUUGU CUGAUGAG X CGAA AAAAAUAA	524	TTATTTT T ACAATGGC	1052
3183	GGCCAUUG CUGAUGAG X CGAA AAAAAUA	525	TATTTT T CAATGGCC	1053
3199	AAAUGGUU CUGAUGAG X CGAA AUUCCAUG	526	CATGGAAT A AACCATTT	1054
3206	UUUGUAAA CUGAUGAG X CGAA AUGGUUUA	527	TAAACCAT T TTTACAAA	1055
3207	UUUUGUAA CUGAUGAG X CGAA AAUGGUUU	528	AAACCATT T TTACAAA	1056

Input Sequence = PTPN1 (Homo sapiens protein tyrosine phosphatase, non-receptor type 1 (PTPN1)
3215 bp)

Cut Site = UH.

--	--	--	--

Table 4

Table 4: Human PTP-1B NCH Ribozyme and Target Sequence

Nt. Position	Ribozyme Sequence	Seq. ID Nos.	Substrate Sequence	Seq. ID Nos.
13	CCGCUCUA CUGAUGAG X CGAA ICCGCGUC	1057	GACGCGGC C TAGAGCGG	1781
14	GCCGCUCU CUGAUGAG X CGAA ICGCGCU	1058	ACGCGGCC T AGAGCGGC	1782
23	GCGCCGUC CUGAUGAG X CGAA ICCGUCU	1059	AGAGCGGC A GACGCGC	1783
32	CGGCCAC CUGAUGAG X CGAA ICGCGUC	1060	GACGCGC A GTGGCCG	1784
39	UCCUUCUC CUGAUGAG X CGAA ICCACUG	1061	CAGTGGC C GAGAAGGA	1785
53	GCGGCUGC CUGAUGAG X CGAA ICGCCUC	1062	GGAGGCGC A GCAGCCG	1786
56	AGGGCGGC CUGAUGAG X CGAA ICUGCGC	1063	GGCGCAGC A GCCGCCCT	1787
59	GCCAGGGC CUGAUGAG X CGAA ICUGCUG	1064	GCAGCAGC C GCCCTGGC	1788
62	CGGGCCAG CUGAUGAG X CGAA ICGCGUC	1065	GCAGCCG C CTGGCCG	1789
63	ACGGGCCA CUGAUGAG X CGAA ICGCGUC	1066	CAGCCGCC C TGGCCCT	1790
64	GACGGGCC CUGAUGAG X CGAA IGGCGCU	1067	AGCCGCC T GGCCCTC	1791
68	CCAUGACG CUGAUGAG X CGAA ICCAGGGC	1068	GCCCTGGC C CGTCATGG	1792
69	UCCAUGAC CUGAUGAG X CGAA IGCAGGG	1069	CCCTGGCC C GTCATGGA	1793
73	CAUCUCCA CUGAUGAG X CGAA IACGGGCC	1070	GGCCGTC A TGGAGATG	1794
98	UGUCGAUC CUGAUGAG X CGAA ICUGAAC	1071	GTTCGAGC A GATCGACA	1795
106	CCCGGACU CUGAUGAG X CGAA IUUGAUCU	1072	AGATCGAC A AGTCCGGG	1796
111	CAGCUCU CUGAUGAG X CGAA IACUUGUC	1073	GACAAGTC C GGGAGCTG	1797
118	GGCGGCC CUGAUGAG X CGAA ICUCGGG	1074	CCGGGAGC T GGGCGGCC	1798
126	UGGUAUU CUGAUGAG X CGAA ICCGCCA	1075	TGGGCGGC C ATTTACCA	1799
127	CUGGUAUU CUGAUGAG X CGAA ICGGCC	1076	GGGCGGC A TTTACCAG	1800
133	GAUAUCCU CUGAUGAG X CGAA IUAAUUGG	1077	CCATTAC C AGGATATC	1801
134	GGUAUCC CUGAUGAG X CGAA IGUAUUG	1078	CATTACC A GGATATCC	1802
142	UUAUGUC CUGAUGAG X CGAA IUAUCCU	1079	AGGATATC C GACATGAA	1803
146	UGGUUCA CUGAUGAG X CGAA IUUGUAU	1080	TATCCGAC A TGAAGCCA	1804
153	AAGUCACU CUGAUGAG X CGAA ICUUAUG	1081	CATGAAGC C AGTGACTT	1805
154	GAAGUCAC CUGAUGAG X CGAA IGUUAU	1082	ATGAAGCC A GTGACTTC	1806
160	ACAUGGGA CUGAUGAG X CGAA IUCACUGG	1083	CCAGTGAC T TCCCATGT	1807
163	UCUACAUG CUGAUGAG X CGAA IAAGUCAC	1084	GTGACTTC C CATGTAGA	1808
164	CUCUACAU CUGAUGAG X CGAA IGAAGUCA	1085	TGACTTCC C ATGTAGAG	1809
165	ACUCUACA CUGAUGAG X CGAA IGGAAGUC	1086	GACTTCCC A TGTAGAGT	1810
177	GGAAGCUU CUGAUGAG X CGAA ICCACUCU	1087	AGAGTGGC C AAGCTTCC	1811
178	AGGAAGCU CUGAUGAG X CGAA IGCCACUC	1088	GAGTGGCC A AGCTTCCT	1812
182	UCUAGGA CUGAUGAG X CGAA ICUUGGCC	1089	GGCCAAGC T TCCTAAGA	1813
185	UGUUCUUA CUGAUGAG X CGAA IAGCUUG	1090	CAAGCTTC C TAAGAACA	1814
186	UUGUUCU CUGAUGAG X CGAA IGAAGCUU	1091	AAGCTTCC T AAGAACA	1815
193	UCGGUUU CUGAUGAG X CGAA IUUCUAG	1092	CTAAGAAC A AAAACCGA	1816
199	CCUAUUUC CUGAUGAG X CGAA IUUUUUGU	1093	ACAAAAAC C GAAATAGG	1817
211	GACGUCUC CUGAUGAG X CGAA IUACCUAU	1094	ATAGGTAC A GAGACGTC	1818
220	AAAGGGAC CUGAUGAG X CGAA IACGUCUC	1095	GAGACGTC A GTCCCTTT	1819
224	GGUCAAAG CUGAUGAG X CGAA IACGACG	1096	CGTCAGTC C CTTTGACC	1820
225	UGGUCAA CUGAUGAG X CGAA IGACUGAC	1097	GTCAGTCC C TTTGACCA	1821
226	AUGGUCAA CUGAUGAG X CGAA IGGACUGA	1098	TCAGTCCC T TTGACCAT	1822
232	CCGACUAU CUGAUGAG X CGAA IUCAAAGG	1099	CCTTTGAC C ATAGTCGG	1823

Table 4

233	UCCGACUA CUGAUGAG X CGAA IGUCAAAG	1100	CTTGACC A TAGTCGGA	1824
248	CUUGAUGU CUGAUGAG X CGAA IUUUAUUC	1101	GATTAAAC T ACATCAAG	1825
251	CUUCUUGA CUGAUGAG X CGAA IUAGUUUA	1102	TAAACTAC A TCAAGAAG	1826
254	UAUCUUCU CUGAUGAG X CGAA IAUGUAGU	1103	ACTACATC A AGAAGATA	1827
268	GUUGAUUA CUGAUGAG X CGAA IUCAUUUA	1104	ATAATGAC T ATATCAAC	1828
274	ACUAGCGU CUGAUGAG X CGAA IAUUAGU	1105	ACTATATC A ACGCTAGT	1829
279	AUCAAACU CUGAUGAG X CGAA ICGUUGAU	1106	ATCAACGC T AGTTTGAT	1830
303	CUCCUUUG CUGAUGAG X CGAA ICUUCUUC	1107	GAAGAAGC C CAAAGGAG	1831
304	ACUCCUUU CUGAUGAG X CGAA IGCUUCUU	1108	AAGAAGCC C AAAGGAGT	1832
305	AACUCCUU CUGAUGAG X CGAA IGGCUUCU	1109	AGAAGCCC A AAGGAGTT	1833
316	GGUAAGAA CUGAUGAG X CGAA IUAACUCC	1110	GGAGTTAC A TTCTTACC	1834
320	CCUGGGUA CUGAUGAG X CGAA IAAUGUAA	1111	TTACATT C TACCCAGG	1835
324	GGGCCUG CUGAUGAG X CGAA IUAAGAAU	1112	ATTCTTAC C CAGGGCCC	1836
325	AGGGCCCU CUGAUGAG X CGAA IGUAAGAA	1113	TTCTTACC C AGGGCCCT	1837
326	AAGGGCCC CUGAUGAG X CGAA IGGUAAGA	1114	TCTTACCC A GGGCCCTT	1838
331	AGGCAAAG CUGAUGAG X CGAA ICCUGGG	1115	CCCAGGGC C CTTGCTT	1839
332	UAGGCAA CUGAUGAG X CGAA IGCCUGG	1116	CCAGGGCC C TTTGCTTA	1840
333	UUAGGCAA CUGAUGAG X CGAA IGGCCUG	1117	CAGGGCCC T TTGCTTA	1841
338	AUGUGUUA CUGAUGAG X CGAA ICAAAGGG	1118	CCCTTTGC C TAACACAT	1842
339	CAUGUGUU CUGAUGAG X CGAA IGCAAAGG	1119	CCTTTGCC T AACACATG	1843
343	ACCGCAUG CUGAUGAG X CGAA IUUAGGCA	1120	TGCCTAAC A CATGCGGT	1844
345	UGACCGCA CUGAUGAG X CGAA IUGUUAGG	1121	CCTAACAC A TGCGGTCA	1845
353	CCCAAAAG CUGAUGAG X CGAA IACCGCAU	1122	ATGCGGTC A CTTTGGG	1846
355	CUCCAAA CUGAUGAG X CGAA IUGACCGC	1123	GCGGTCAC T TTTGGGAG	1847
377	UGCUIUUC CUGAUGAG X CGAA ICUCCAC	1124	GTGGGAGC A GAAAAGCA	1848
385	GACACCCC CUGAUGAG X CGAA ICUUUCU	1125	AGAAAAGC A GGGGTGTC	1849
397	GUUGAGCA CUGAUGAG X CGAA IACGACAC	1126	GTGTCGTC A TGCTCAAC	1850
401	CUCUGUUG CUGAUGAG X CGAA ICAUGACG	1127	CGTCATGC T CAACAGAG	1851
403	CACUCUGU CUGAUGAG X CGAA IAGCAUGA	1128	TCATGCTC A ACAGAGTG	1852
406	CAUCACUC CUGAUGAG X CGAA IUUGAGCA	1129	TGCTCAAC A GAGTGATG	1853
438	CAGUAUUG CUGAUGAG X CGAA ICGCAUUU	1130	AAATGCGC A CAATACTG	1854
440	GCCAGUAU CUGAUGAG X CGAA IUGCGCAU	1131	ATGCGCAC A ATACTGGC	1855
445	UUGUGGCC CUGAUGAG X CGAA IUUAUUGU	1132	CACAATAC T GGCCACAA	1856
449	CUUUUUGU CUGAUGAG X CGAA ICCAGUAU	1133	ATACTGGC C AAAAAAG	1857
450	UCUUUUUG CUGAUGAG X CGAA IGCCAGUA	1134	TACTGGCC A CAAAAAGA	1858
452	CUUCUUUU CUGAUGAG X CGAA IUGGCCAG	1135	CTGGCCAC A AAAAGAAG	1859
475	GUCUUCAA CUGAUGAG X CGAA IAUCAUCU	1136	AGATGATC T TTGAAGAC	1860
484	CAAAUUUG CUGAUGAG X CGAA IUCUUCAA	1137	TTGAAGAC A CAAATTTG	1861
486	UUCAAAUU CUGAUGAG X CGAA IUGUCUUC	1138	GAAGACAC A AATTTGAA	1862
501	GAGAUCAA CUGAUGAG X CGAA IUUAAUUU	1139	AAATTAAC A TTGATCTC	1863
508	AUCUUCAG CUGAUGAG X CGAA IAUCAAUG	1140	CATTGATC T CTGAAGAT	1864
510	AUAUCUUC CUGAUGAG X CGAA IAGAUCAA	1141	TTGATCTC T GAAGATAT	1865
520	AUAUGACU CUGAUGAG X CGAA IAUUCUU	1142	AAGATATC A AGTCATAT	1866
525	GUUAAUUA CUGAUGAG X CGAA IACUUGAU	1143	ATCAAGTC A TATTATAC	1867
534	UGUCGCAC CUGAUGAG X CGAA IUUAUUA	1144	TATTATAC A GTGCGACA	1868
542	AUUCUAGC CUGAUGAG X CGAA IUCGCACU	1145	AGTGCAC A GCTAGAAT	1869
545	CCAAUUCU CUGAUGAG X CGAA ICUGUCGC	1146	GCGACAGC T AGAATTGG	1870

Table 4

559	GGUUGUAA CUGAUGAG X CGAA IUUUUCCA	1147	TGGAAAAC C TTACAACC	1871
560	GGGUGUGA CUGAUGAG X CGAA IGUUUUCC	1148	GGAAAACC T TACAACCC	1872
564	UCUUGGGU CUGAUGAG X CGAA IUAAGGUU	1149	AACCTTAC A ACCCAAGA	1873
567	GUUUCUUG CUGAUGAG X CGAA IUUGUAAG	1150	CTTACAAC C CAAGAAAC	1874
568	AGUUUCUU CUGAUGAG X CGAA IGUUGUAA	1151	TTACAACC C AAGAAACT	1875
569	GAGUUUCU CUGAUGAG X CGAA IGGUUGUA	1152	TACAACCC A AGAAACTC	1876
576	AUCUCUCG CUGAUGAG X CGAA IUUUCUUG	1153	CAAGAAAC T CGAGAGAT	1877
586	GAA AUGUA CUGAUGAG X CGAA IAUCUCUC	1154	GAGAGATC T TACATTTC	1878
590	AGUGGAAA CUGAUGAG X CGAA IUAAGAUC	1155	GATCTTAC A TTTCCACT	1879
595	GGUAUAGU CUGAUGAG X CGAA IAAAUGUA	1156	TACATTTC C ACTATACC	1880
596	UGGUAUAG CUGAUGAG X CGAA IGAAAUGU	1157	ACATTTC C A CTATACCA	1881
598	UGUGGUAU CUGAUGAG X CGAA IUGGAAAU	1158	ATTTCAC T ATACCACA	1882
603	GGCCAUGU CUGAUGAG X CGAA IUUAUGUG	1159	CACTATAC C ACATGGCC	1883
604	AGGCCAUG CUGAUGAG X CGAA IGUAUAGU	1160	ACTATACC A CATGGCCT	1884
606	UCAGGCCA CUGAUGAG X CGAA IUGGUAVA	1161	TATACCAC A TGGCCTGA	1885
611	CAAAGUCA CUGAUGAG X CGAA ICCAUGUG	1162	CACATGGC C TGACTTTG	1886
612	CCAAAGUC CUGAUGAG X CGAA IGCCAUGU	1163	ACATGGCC T GACTTTGG	1887
616	GACUCCAA CUGAUGAG X CGAA IUCAGGCC	1164	GGCCTGAC T TTGGAGTC	1888
625	UGAUUCAG CUGAUGAG X CGAA IACUCCAA	1165	TTGGAGTC C CTGAATCA	1889
626	GUGAUUCA CUGAUGAG X CGAA IGACUCCA	1166	TGGAGTCC C TGAATCAC	1890
627	GGUGAUUC CUGAUGAG X CGAA IGGACUCC	1167	GGAGTCCC T GAATCACC	1891
633	GAGGCUGG CUGAUGAG X CGAA IAUUCAGG	1168	CCTGAATC A CCAGCCTC	1892
635	AUGAGGCU CUGAUGAG X CGAA IUGAUUCA	1169	TGAATCAC C AGCCTCAT	1893
636	AAUGAGGC CUGAUGAG X CGAA IGUGAUUC	1170	GAATCACC A GCCTCATT	1894
639	AAGAAUGA CUGAUGAG X CGAA ICUGGUGA	1171	TCACCAGC C TCATTCTT	1895
640	CAAGAAUG CUGAUGAG X CGAA IGCUGGUG	1172	CACCAGCC T CATTCTTG	1896
642	UUAAGAA CUGAUGAG X CGAA IAGGCUGG	1173	CCAGCCTC A TTCTTGAA	1897
646	AAAGUUCA CUGAUGAG X CGAA IAAUGAGG	1174	CCTCATTC T TGAAC TTT	1898
652	GAAAAGAA CUGAUGAG X CGAA IUUCAAGA	1175	TCTTGAAC T TTCTTTTC	1899
656	CUUUGAAA CUGAUGAG X CGAA IAAAGUUC	1176	GAAC TTT TTTCAAAG	1900
661	UCGACUU CUGAUGAG X CGAA IAAAAGAA	1177	TTCTTTTC A AAGTCCGA	1901
667	UGACUCUC CUGAUGAG X CGAA IACUUUGA	1178	TCAAAGTC C GAGAGTCA	1902
675	AGUGACCC CUGAUGAG X CGAA IACUCUCG	1179	CGAGAGTC A GGGTCACT	1903
681	GGGUCUG CUGAUGAG X CGAA IACCCUGA	1180	TCAGGGTC A CTCAGCCC	1904
683	CCGGGUCG CUGAUGAG X CGAA IUGACCCU	1181	AGGGTCAC T CAGCCGG	1905
685	CUCGGGCG CUGAUGAG X CGAA IAGUGACC	1182	GGTCACTC A GCCCGGAG	1906
688	GUGCUCCG CUGAUGAG X CGAA ICUGAGUG	1183	CACTCAGC C CGGAGCAC	1907
689	CGUGCUC CUGAUGAG X CGAA IGCUGAGU	1184	ACTCAGCC C GGAGCACG	1908
695	CGGGCCCG CUGAUGAG X CGAA ICUCGGGG	1185	CCCGGAGC A CGGGCCCG	1909
701	CCACAACG CUGAUGAG X CGAA ICCCGUGC	1186	GCACGGGC C CGTTGTGG	1910
702	ACCACAAC CUGAUGAG X CGAA IGCCCGUG	1187	CACGGGCC C GTTGTGGT	1911
713	CACUGCAG CUGAUGAG X CGAA ICACCACA	1188	TGTGTGTC A CTGCAGTG	1912
715	UGCACUGC CUGAUGAG X CGAA IUGCACCA	1189	TGGTGCAC T GCAGTGCA	1913
718	GCCUGCAC CUGAUGAG X CGAA ICAGUGCA	1190	TGCACTGC A GTGCAGGC	1914
723	CCGAUGCC CUGAUGAG X CGAA ICACUGCA	1191	TGCACTGC A GGCATCGG	1915
727	CCUGCCGA CUGAUGAG X CGAA ICCUGCAC	1192	GTGCAGGC A TCGGCAGG	1916
733	UCCAGACC CUGAUGAG X CGAA ICCGAUGC	1193	GCATCGGC A GGTCTGGA	1917

Table 4

738	AAGGUUCC CUGAUGAG X CGAA IACCUGCC	1194	GGCAGGTC T GGAACCTT	1918
744	AGACAGAA CUGAUGAG X CGAA IUUCCAGA	1195	TCTGGAAC C TTCTGTCT	1919
745	CAGACAGA CUGAUGAG X CGAA IGUUCCAG	1196	CTGGAACC T TCTGTCTG	1920
748	AGCCAGAC CUGAUGAG X CGAA IAAGGUUC	1197	GAACCTTC T GTCTGGCT	1921
752	UAUCAGCC CUGAUGAG X CGAA IACAGAAG	1198	CTTCTGTC T GGCTGATA	1922
756	CAGGUUUC CUGAUGAG X CGAA ICCAGACA	1199	TGTCTGGC T GATACCTG	1923
762	AAGAGGCA CUGAUGAG X CGAA IUAUCAGC	1200	GCTGATAC C TGCCTCTT	1924
763	CAAGAGGC CUGAUGAG X CGAA IGUAUCAG	1201	CTGATACC T GCCTCTTG	1925
766	CAGCAAGA CUGAUGAG X CGAA ICAGGAU	1202	ATACCTGC C TCTTGCTG	1926
767	UCAGCAAG CUGAUGAG X CGAA IGCAGGUA	1203	TACCTGCC T CTTGCTGA	1927
769	CAUCAGCA CUGAUGAG X CGAA IAGGCAGG	1204	CCTGCCTC T TGCTGATG	1928
773	UGUCCAUC CUGAUGAG X CGAA ICAAGAGG	1205	CCTCTTGC T GATGGACA	1929
781	UUUCCUCU CUGAUGAG X CGAA IUCCAUCA	1206	TGATGGAC A AGAGGAAA	1930
793	GGAAGAAG CUGAUGAG X CGAA IUUUUCC	1207	GGAAAGAC C CTTCTTCC	1931
794	CGGAAGAA CUGAUGAG X CGAA IGUCUUUC	1208	GAAAGACC C TTCTTCCG	1932
795	ACGGAAGA CUGAUGAG X CGAA IGGUCUUU	1209	AAAGACCC T TCTTCCGT	1933
798	UCAACGGA CUGAUGAG X CGAA IAAGGGUC	1210	GACCCTTC T TCCGTTGA	1934
801	AUAUCAAC CUGAUGAG X CGAA IAAGAAGG	1211	CCTTCTTC C GTTGATAT	1935
811	CACUUUCU CUGAUGAG X CGAA IUAUCAA	1212	TTGATATC A AGAAAGTG	1936
821	UUUCUAAC CUGAUGAG X CGAA ICACUUUC	1213	GAAAGTGC T GTTAGAAA	1937
851	UCUGGAUC CUGAUGAG X CGAA ICCCAUC	1214	GATGGGGC T GATCCAGA	1938
856	GGCUGUCU CUGAUGAG X CGAA IAUACGCC	1215	GGCTGATC C AGACAGCC	1939
857	CGCUGUC CUGAUGAG X CGAA IGAUCAGC	1216	GCTGATCC A GACAGCCG	1940
861	UGGUCGGC CUGAUGAG X CGAA IUCUGGAU	1217	ATCCAGAC A GCCGACCA	1941
864	AGCUGGUC CUGAUGAG X CGAA ICUGUCUG	1218	CAGACAGC C GACCAGCT	1942
868	GCGCAGCU CUGAUGAG X CGAA IUCGGCUG	1219	CAGCCGAC C AGCTGCGC	1943
869	AGCGCAGC CUGAUGAG X CGAA IGUCGGCU	1220	AGCCGACC A GCTGCGCT	1944
872	AGAAGCGC CUGAUGAG X CGAA ICUGGUCG	1221	CGACCAGC T GCGTTCT	1945
877	GUAGGAGA CUGAUGAG X CGAA ICGCAGCU	1222	AGCTGCGC T TCTCCTAC	1946
880	CAGGUAGG CUGAUGAG X CGAA IAAGCGCA	1223	TGCGCTTC T CCTACCTG	1947
882	GCCAGGUA CUGAUGAG X CGAA IAGAAGCG	1224	CGCTTCTC C TACCTGGC	1948
883	AGCCAGGU CUGAUGAG X CGAA IGAGAAGC	1225	GCTTCTCC T ACCTGGCT	1949
886	CACAGCCA CUGAUGAG X CGAA IUAGGAGA	1226	TCTCCTAC C TGGCTGTG	1950
887	UCACAGCC CUGAUGAG X CGAA IGUAGGAG	1227	CTCCTACC T GGCTGTGA	1951
891	UCGAUCAC CUGAUGAG X CGAA ICCAGGUA	1228	TACCTGGC T GTGATCGA	1952
906	AUGAAUUU CUGAUGAG X CGAA ICACCUUC	1229	GAAGGTGC C AAATTCAT	1953
907	GAUGAAUU CUGAUGAG X CGAA IGCACCUU	1230	AAGGTGCC A AATTCATC	1954
913	CCCAUGA CUGAUGAG X CGAA IAAUUUGG	1231	CCAAATTC A TCATGGGG	1955
916	GUCCCCCA CUGAUGAG X CGAA IAUGAAUU	1232	AATTCATC A TGGGGGAC	1956
925	CACGGAAG CUGAUGAG X CGAA IUCCCCCA	1233	TGGGGGAC T CTTCCGTG	1957
927	UGCACGGA CUGAUGAG X CGAA IAGUCCCC	1234	GGGGACTC T TCCGTGCA	1958
930	UCCUGCAC CUGAUGAG X CGAA IAAGAGUC	1235	GACTCTTC C GTGCAGGA	1959
935	ACUGAUCC CUGAUGAG X CGAA ICACGGAA	1236	TTCCGTGC A GGATCAGT	1960
941	CCUUCAC CUGAUGAG X CGAA IAUCCUGC	1237	GCAGGATC A GTGGAAGG	1961
953	CGUGGGAA CUGAUGAG X CGAA ICUCUUC	1238	GAAGGAGC T TTCCACG	1962
957	UCCUCGUG CUGAUGAG X CGAA IAAAGCUC	1239	GAGCTTTC C CACGAGGA	1963
958	GUCCUCGU CUGAUGAG X CGAA IGAAAGCU	1240	AGCTTTCC C ACGAGGAC	1964

Table 4

959	GGUCCUCG CUGAUGAG X CGAA IGGAAAGC	1241	GCTTTCCC A CGAGGACC	1965
967	GGGCUCCA CUGAUGAG X CGAA IUCCUCGU	1242	ACGAGGAC C TGGAGCCC	1966
968	GGGGCUCC CUGAUGAG X CGAA IGUCCUCG	1243	CGAGGACC T GGAGCCCC	1967
974	CGGGUGGG CUGAUGAG X CGAA ICUCCAGG	1244	CCTGGAGC C CCCACCCG	1968
975	UCGGGUGG CUGAUGAG X CGAA IGUCCAG	1245	CTGGAGCC C CCACCCGA	1969
976	CUCGGGUG CUGAUGAG X CGAA IGGCUCCA	1246	TGGAGCCC C CACCCGAG	1970
977	GCUCGGGU CUGAUGAG X CGAA IGGGCUCC	1247	GGAGCCCC C ACCCGAGC	1971
978	UGCUCGGG CUGAUGAG X CGAA IGGGGCUC	1248	GAGCCCCC A CCCGAGCA	1972
980	UAUGCUCG CUGAUGAG X CGAA IUGGGGGC	1249	GCCCCCAC C CGAGCATA	1973
981	AUAUGCUC CUGAUGAG X CGAA IGUGGGGG	1250	CCCCCACC C GAGCATAT	1974
986	GGGGGAUA CUGAUGAG X CGAA ICUCGGGU	1251	ACCCGAGC A TATCCCCC	1975
991	AGGUGGGG CUGAUGAG X CGAA IAUAUGCU	1252	AGCATATC C CCCCACCT	1976
992	GAGGUGGG CUGAUGAG X CGAA IGAUAUGC	1253	GCATATCC C CCCACCTC	1977
993	GGAGGUGG CUGAUGAG X CGAA IGGAUUAG	1254	CATATCCC C CCACCTCC	1978
994	GGGAGGUG CUGAUGAG X CGAA IGGGAUUA	1255	ATATCCCC C CACCTCCC	1979
995	GGGGAGGU CUGAUGAG X CGAA IGGGGAUA	1256	TATCCCCC C ACCTCCCC	1980
996	CGGGGAGG CUGAUGAG X CGAA IGGGGGAU	1257	ATCCCCC A CCTCCCCG	1981
998	GCCGGGGA CUGAUGAG X CGAA IUGGGGGG	1258	CCCCCAC C TCCCCGGC	1982
999	GGCCGGGG CUGAUGAG X CGAA IGUGGGGG	1259	CCCCCACC T CCCCCGCC	1983
1001	GUGGCCGG CUGAUGAG X CGAA IAGGUGGG	1260	CCCACCTC C CCGGCCAC	1984
1002	GGUGGCCG CUGAUGAG X CGAA IGAGGUGG	1261	CCACCTCC C CGGCCACC	1985
1003	GGGUGGCC CUGAUGAG X CGAA IGGAGGUG	1262	CACCTCCC C GGCCACCC	1986
1007	GUUUGGGU CUGAUGAG X CGAA ICCGGGGA	1263	TCCCCGGC C ACCCAAAC	1987
1008	CGUUGGG CUGAUGAG X CGAA IGCCGGGG	1264	CCCCGGCC A CCCAAACG	1988
1010	UUCGUUUG CUGAUGAG X CGAA IUGGCCGG	1265	CCGGCCAC C CAAACGAA	1989
1011	AUUCGUUU CUGAUGAG X CGAA IGUGGCCG	1266	CGGCCACC C AAACGAAT	1990
1012	GAUUCGUU CUGAUGAG X CGAA IGGUGGCC	1267	GGCCACCC A AACGAATC	1991
1021	UGGCUCCA CUGAUGAG X CGAA IAUUCGUU	1268	AACGAATC C TGGAGCCA	1992
1022	GUGGCUCC CUGAUGAG X CGAA IGAUUCGU	1269	ACGAATCC T GGAGCCAC	1993
1028	CAUUGUGU CUGAUGAG X CGAA ICUCCAGG	1270	CCTGGAGC C ACACAATG	1994
1029	CCAUUGUG CUGAUGAG X CGAA IGUCCAG	1271	CTGGAGCC A CACAATGG	1995
1031	UCCCAUUG CUGAUGAG X CGAA IUGGCUCC	1272	GGAGCCAC A CAATGGGA	1996
1033	UUUCCCAU CUGAUGAG X CGAA IUUGGCUU	1273	AGCCACAC A ATGGGAAA	1997
1045	GAACUCCC CUGAUGAG X CGAA ICAUUUCC	1274	GGAAATGC A GGGAGTTC	1998
1054	AUUUGGGA CUGAUGAG X CGAA IACUCCC	1275	GGGAGTTC T TCCCAAAT	1999
1057	GUGAUUUG CUGAUGAG X CGAA IAAGAACU	1276	AGTTCTTC C CAAATCAC	2000
1058	GGUGAUUU CUGAUGAG X CGAA IGAAGAAC	1277	GTTCTTCC C AAATCACC	2001
1059	UGGUGAUU CUGAUGAG X CGAA IGAAGAA	1278	TTCTTCCC A AATCACCA	2002
1064	CCCACUGG CUGAUGAG X CGAA IAUUUGGG	1279	CCCCAAATC A CCAGTGGG	2003
1066	CACCCACU CUGAUGAG X CGAA IUGAUUUG	1280	CAAATCAC C AGTGGGTG	2004
1067	UCACCCAC CUGAUGAG X CGAA IGUGAUUU	1281	AAATCAC A GTGGGTGA	2005
1086	UCCUCCUG CUGAUGAG X CGAA IUCUCUUC	1282	GAAGAGAC C CAGGAGGA	2006
1087	AUCCUCCU CUGAUGAG X CGAA IGUCUCUU	1283	AAGAGACC C AGGAGGAT	2007
1088	UAUCCUCC CUGAUGAG X CGAA IGGUCUCU	1284	AGAGACCC A GGAGGATA	2008
1102	GAUGGGGC CUGAUGAG X CGAA IUCUUUAU	1285	ATAAAGAC T GCCCCATC	2009
1105	CUUGAUGG CUGAUGAG X CGAA ICAGUCUU	1286	AAGACTGC C CCATCAAG	2010
1106	CCUUGAUG CUGAUGAG X CGAA IGCAGUCU	1287	AGACTGCC C CATCAAGG	2011

Table 4

1107	UCCUUGAU CUGAUGAG X CGAA IGGCAGUC	1288	GACTGCCC C ATCAAGGA	2012
1108	UUCUUGA CUGAUGAG X CGAA IGGCAGU	1289	ACTGCCCC A TCAAGGAA	2013
1111	UUCUUCU CUGAUGAG X CGAA IAUGGGG	1290	GCCCCATC A AGGAAGAA	2014
1129	AUUUAAGG CUGAUGAG X CGAA ICUUCCU	1291	AAGGAAGC C CCTTAAAT	2015
1130	CAUUUAAG CUGAUGAG X CGAA IGCUCCU	1292	AGGAAGCC C CTTAAATG	2016
1131	GCAUUUA CUGAUGAG X CGAA IGGCUCC	1293	GGAAGCCC C TTAAATGC	2017
1132	GGCAUUUA CUGAUGAG X CGAA IGGCUUC	1294	GAAGCCCC T TAAATGCC	2018
1140	UAGGGUGC CUGAUGAG X CGAA ICAUUUA	1295	TTAAATGC C GCACCCTA	2019
1143	CCGUAGGG CUGAUGAG X CGAA ICGCAUU	1296	AATGCCGC A CCCTACGG	2020
1145	UGCCGUAG CUGAUGAG X CGAA IUGCGGC	1297	TGCCGCAC C CTACGGCA	2021
1146	AUGCCGUA CUGAUGAG X CGAA IGUGCGG	1298	GCCGCACC C TACGGCAT	2022
1147	GAUGCCGU CUGAUGAG X CGAA IGGUGCG	1299	CCGCACCC T ACGGCATC	2023
1153	GCUUUCGA CUGAUGAG X CGAA ICCGUAGG	1300	CCTACGGC A TCGAAAGC	2024
1162	UUGACUCA CUGAUGAG X CGAA ICUUUCGA	1301	TCGAAAGC A TGAGTCAA	2025
1169	CAGUGUCU CUGAUGAG X CGAA IACUCAUG	1302	CATGAGTC A AGACACTG	2026
1174	AACUUCAG CUGAUGAG X CGAA IUUCUGAC	1303	GTCAAGAC A CTGAAGTT	2027
1176	CUAACUUC CUGAUGAG X CGAA IUGUCUUG	1304	CAAGACAC T GAAGTTAG	2028
1208	CACCUCGA CUGAUGAG X CGAA IACUCCC	1305	GGGAAGTC T TCGAGGTG	2029
1218	GCAGCCUG CUGAUGAG X CGAA ICACCUCG	1306	CGAGGTGC C CAGGCTGC	2030
1219	GGCAGCCU CUGAUGAG X CGAA IGCACCUC	1307	GAGGTGCC C AGGCTGCC	2031
1220	AGGCAGCC CUGAUGAG X CGAA IGGACCU	1308	AGGTGCCC A GGCTGCCT	2032
1224	GGGAGGC CUGAUGAG X CGAA ICCUGGG	1309	GCCCAGGC T GCCTCCCC	2033
1227	GCUGGGGA CUGAUGAG X CGAA ICAGCCUG	1310	CAGGCTGC C TCCCCAGC	2034
1228	GGCUGGG CUGAUGAG X CGAA ICAGCCU	1311	AGGCTGCC T CCCCAGCC	2035
1230	UUGGCUG CUGAUGAG X CGAA IAGGCAGC	1312	GCTGCCTC C CCAGCCAA	2036
1231	UUUGGCUG CUGAUGAG X CGAA IGAGGCAG	1313	CTGCCTCC C CAGCCAA	2037
1232	CUUUGGC CUGAUGAG X CGAA IGGAGGCA	1314	TGCCTCCC C AGCCAAAG	2038
1233	CCUUGGC CUGAUGAG X CGAA IGGAGGC	1315	GCCTCCCC A GCCAAAGG	2039
1236	UCCCCUU CUGAUGAG X CGAA ICUGGGGA	1316	TCCCCAGC C AAAGGGGA	2040
1237	CUCCCCU CUGAUGAG X CGAA ICUGGGG	1317	CCCCAGCC A AAGGGGAG	2041
1247	GCAGUGAC CUGAUGAG X CGAA ICUCCCCU	1318	AGGGGAGC C GTCAGTGC	2042
1251	UCGGGCAG CUGAUGAG X CGAA IACGGCUC	1319	GAGCCGTC A CTGCCGA	2043
1253	UCUCGGG CUGAUGAG X CGAA IUGACGGC	1320	GCCGTCAC T GCCCGAGA	2044
1256	CCUUCUC CUGAUGAG X CGAA ICAGUGAC	1321	GTCAGTGC C CGAGAAGG	2045
1257	UCCUUCUC CUGAUGAG X CGAA ICAGUGA	1322	TCACTGCC C GAGAAGGA	2046
1273	CAGUGCAU CUGAUGAG X CGAA IUCCUCGU	1323	ACGAGGAC C ATGCACTG	2047
1274	UCAGUGCA CUGAUGAG X CGAA IGUCCUCG	1324	CGAGGACC A TGCACTGA	2048
1278	UAACUCAG CUGAUGAG X CGAA ICAUGGUC	1325	GACCATGC A CTGAGTTA	2049
1280	AGUAACUC CUGAUGAG X CGAA IUGCAUGG	1326	CCATGCAC T GAGTTACT	2050
1288	GGGCUUC CUGAUGAG X CGAA IUAACUCA	1327	TGAGTTAC T GGAAGCCC	2051
1295	CCAGGAAG CUGAUGAG X CGAA ICUCCAG	1328	CTGGAAGC C CTCCTGG	2052
1296	ACCAGGAA CUGAUGAG X CGAA IGCUCCA	1329	TGGAAGCC C TTCCTGGT	2053
1297	GACCAGGA CUGAUGAG X CGAA IGGCUUC	1330	GGAAGCCC T TCCTGGTC	2054
1300	GUUGACCA CUGAUGAG X CGAA IAAGGGCU	1331	AGCCCTTC C TGGTCAAC	2055
1301	UGUGACC CUGAUGAG X CGAA IGAAGGGC	1332	GCCCTTCC T GGTCAACA	2056
1306	GCACAUG CUGAUGAG X CGAA IACCAGGA	1333	TCCTGGTC A ACATGTGC	2057
1309	CACGCACA CUGAUGAG X CGAA IUUGACCA	1334	TGGTCAAC A TGTGCGTG	2058

Table 4

1320	AGGACCGU CUGAUGAG X CGAA ICCACGCA	1335	TGCGTGGC T ACGGTCCT	2059
1327	GGCCGUGA CUGAUGAG X CGAA IACCGUAG	1336	CTACGGTC C TCACGGCC	2060
1328	CGGCCGUG CUGAUGAG X CGAA IGACCGUA	1337	TACGGTCC T CACGGCCG	2061
1330	GCCGGCCG CUGAUGAG X CGAA IAGGACCG	1338	CGGTCTC A CGGCCGGC	2062
1335	UAAGCGCC CUGAUGAG X CGAA ICCGUGAG	1339	CTCACGGC C GCGCTTA	2063
1341	CAGAGGUA CUGAUGAG X CGAA ICGCCGGC	1340	GCCGGCGC T TACCTCTG	2064
1345	GUAGCAGA CUGAUGAG X CGAA IUAAGCGC	1341	GCGCTTAC C TCTGCTAC	2065
1346	UGUAGCAG CUGAUGAG X CGAA IGUAAGCG	1342	CGCTTACC T CTGCTACA	2066
1348	CCUGUAGC CUGAUGAG X CGAA IAGGUAAG	1343	CTTACCTC T GCTACAGG	2067
1351	GAACCUGU CUGAUGAG X CGAA ICAGAGGU	1344	ACCTCTGC T ACAGGTTC	2068
1354	CAGGAACC CUGAUGAG X CGAA IUAGCAGA	1345	TCTGCTAC A GGTTCCTG	2069
1360	GUUGAACA CUGAUGAG X CGAA IAACCUGU	1346	ACAGGTTC C TGTTC AAC	2070
1361	UGUGAAC CUGAUGAG X CGAA IGAACCUG	1347	CAGGTTC T GTTCAACA	2071
1366	GUUGCUGU CUGAUGAG X CGAA IACAGGA	1348	TCCTGTTC A ACAGCAAC	2072
1369	UGUGUUGC CUGAUGAG X CGAA IUUGAACA	1349	TGTTCAAC A GCAACACA	2073
1372	CUAUGUGU CUGAUGAG X CGAA ICUGUUGA	1350	TCAACAGC A ACACATAG	2074
1375	AGGCUAUG CUGAUGAG X CGAA IUUGCUGU	1351	ACAGCAAC A CATAGCCT	2075
1377	UCAGGCUA CUGAUGAG X CGAA IUGUUGCU	1352	AGCAACAC A TAGCCTGA	2076
1382	GAGGGUCA CUGAUGAG X CGAA ICUAUGUG	1353	CACATAGC C TGACCTC	2077
1383	GGAGGGUC CUGAUGAG X CGAA IGCUAUGU	1354	ACATAGCC T GACCCTCC	2078
1387	UGGAGGAG CUGAUGAG X CGAA IUCAGGCU	1355	AGCCTGAC C CTCCTCCA	2079
1388	GUGGAGGA CUGAUGAG X CGAA IGUCAGGC	1356	GCCTGACC C TCCTCCAC	2080
1389	AGUGGAGG CUGAUGAG X CGAA IGGUCAGG	1357	CCTGACCC T CCTCCACT	2081
1391	GGAGUGGA CUGAUGAG X CGAA IAGGGUCA	1358	TGACCTC C TCCACTCC	2082
1392	UGGAGUGG CUGAUGAG X CGAA IGAGGGUC	1359	GACCTCC T CCACTCCA	2083
1394	GGUGGAGU CUGAUGAG X CGAA IAGGAGGG	1360	CCCTCCTC C ACTCCACC	2084
1395	AGGUGGAG CUGAUGAG X CGAA IGAGGAGG	1361	CCTCCTCC A CTCCACCT	2085
1397	GGAGGUGG CUGAUGAG X CGAA IUGGAGGA	1362	TCCTCCAC T CCACCTCC	2086
1399	GUGGAGGU CUGAUGAG X CGAA IAGUGGAG	1363	CTCCACTC C ACCTCCAC	2087
1400	GGUGGAGG CUGAUGAG X CGAA IGAGUGGA	1364	TCCACTCC A CCTCCACC	2088
1402	UGGGUGGA CUGAUGAG X CGAA IUGGAGUG	1365	CACTCCAC C TCCACCCA	2089
1403	GUGGGUGG CUGAUGAG X CGAA IGUGGAGU	1366	ACTCCACC T CCACCCAC	2090
1405	CAGUGGGU CUGAUGAG X CGAA IAGGUGGA	1367	TCCACCTC C ACCCACTG	2091
1406	ACAGUGGG CUGAUGAG X CGAA IGAGGUGG	1368	CCACCTCC A CCCACTGT	2092
1408	GGACAGUG CUGAUGAG X CGAA IUGGAGGU	1369	ACCTCCAC C CACTGTCC	2093
1409	CGACAGU CUGAUGAG X CGAA IGUGGAGG	1370	CCTCCACC C ACTGTCCG	2094
1410	GCGGACAG CUGAUGAG X CGAA IGGUGGAG	1371	CTCCACCC A CTGTCCGC	2095
1412	AGGCGGAC CUGAUGAG X CGAA IUGGGUGG	1372	CCACCCAC T GTCCGCCT	2096
1416	GCAGAGGC CUGAUGAG X CGAA IACAGUGG	1373	CCACTGTC C GCCTCTGC	2097
1419	CGGCAGAG CUGAUGAG X CGAA ICGGACAG	1374	CTGTCCGC C TCTGCCCG	2098
1420	GCGGCAG CUGAUGAG X CGAA ICGGACA	1375	TGTCCGCC T CTGCCCGC	2099
1422	CUGCGGC CUGAUGAG X CGAA IAGGCGGA	1376	TCCGCCTC T GCCGCAG	2100
1425	GCUCUGCG CUGAUGAG X CGAA ICAGAGGC	1377	GCCTCTGC C CGCAGAGC	2101
1426	GGCUCUGC CUGAUGAG X CGAA IGCAGAGG	1378	CCTCTGCC C GCAGAGCC	2102
1429	GUGGGCUC CUGAUGAG X CGAA ICGGCAG	1379	CTGCCCGC A GAGCCAC	2103
1434	CGGGCGUG CUGAUGAG X CGAA ICUCUGCG	1380	CGCAGAGC C CACGCCCG	2104
1435	UCGGGCGU CUGAUGAG X CGAA IGCUCUGC	1381	GCAGAGCC C ACGCCCGA	2105

Table 4

1436	GUCGGGCG CUGAUGAG X CGAA IGGCUCUG	1382	CAGAGCCC A CGCCCGAC	2106
1440	GCUAGUCG CUGAUGAG X CGAA ICGUGGGC	1383	GCCCACGC C CGACTAGC	2107
1441	UGCUAGUC CUGAUGAG X CGAA ICGUGGG	1384	CCCACGCC C GACTAGCA	2108
1445	UGCCUGCU CUGAUGAG X CGAA IUCGGGCG	1385	CGCCCGAC T AGCAGGCA	2109
1449	GGCAUGCC CUGAUGAG X CGAA ICUAGUCG	1386	CGACTAGC A GGCATGCC	2110
1453	CCGCGGCA CUGAUGAG X CGAA ICCUGCUA	1387	TAGCAGGC A TGCCGCGG	2111
1457	CCUACCGC CUGAUGAG X CGAA ICAUGCCU	1388	AGGCATGC C GCGGTAGG	2112
1473	GGUCCGGC CUGAUGAG X CGAA ICCCUUAC	1389	GTAAGGGC C GCCGGACC	2113
1476	CGCGGUCC CUGAUGAG X CGAA ICGGCCCC	1390	AGGGCCGC C GGACCGCG	2114
1481	CUCUACGC CUGAUGAG X CGAA IUCGGGCG	1391	CGCCGGAC C GCGTAGAG	2115
1493	CGGGGCCC CUGAUGAG X CGAA ICUCUCUA	1392	TAGAGAGC C GGGCCCCG	2116
1498	CCGUCCGG CUGAUGAG X CGAA ICCCGGCU	1393	AGCCGGGC C CCGGACGG	2117
1499	UCCGUCCG CUGAUGAG X CGAA IGCCCGGC	1394	GCCGGGCC C CGGACGGA	2118
1500	GUCCGUCC CUGAUGAG X CGAA IGGCCCGG	1395	CCGGGCCC C GGACGGAC	2119
1517	UUUAGUGC CUGAUGAG X CGAA IAACCAAC	1396	GTTGGTTC T GCACTAAA	2120
1520	GGUUUUAG CUGAUGAG X CGAA ICAGAACC	1397	GGTTCGTC A CTAACC	2121
1522	UGGUUUU CUGAUGAG X CGAA IUCAGAA	1398	TTCTGCAC T AAAACCCA	2122
1528	GGAAGAUG CUGAUGAG X CGAA IUUUUAGU	1399	ACTAAAC C CATCTTCC	2123
1529	GGGAAGAU CUGAUGAG X CGAA IGUUUUAG	1400	CTAAACC C ATCTTCCC	2124
1530	GGGAAGA CUGAUGAG X CGAA IGUUUUUA	1401	TAAACCC A TCTTCCC	2125
1533	UCCGGGGA CUGAUGAG X CGAA IAUGGGUU	1402	AACCATC T TCCCCGA	2126
1536	ACAUCCGG CUGAUGAG X CGAA IAAGAUGG	1403	CCATCTTC C CCGATGT	2127
1537	CACAUCCG CUGAUGAG X CGAA IGAAGAUG	1404	CATCTTCC C CGGATGTG	2128
1538	ACACAUCC CUGAUGAG X CGAA IGGAAGAU	1405	ATCTTCCC C GGATGTGT	2129
1550	GAGGGGUG CUGAUGAG X CGAA IACACACA	1406	TGTGTGTC T CACCCCTC	2130
1552	AUGAGGGG CUGAUGAG X CGAA IAGACACA	1407	TGTGTCTC A CCCCTCAT	2131
1554	GGAUGAGG CUGAUGAG X CGAA IUGAGACA	1408	TGTCTCAC C CCTCATCC	2132
1555	AGGAUGAG CUGAUGAG X CGAA IGUGAGAC	1409	GTCTCACC C CTCATCCT	2133
1556	AAGGAUGA CUGAUGAG X CGAA IGGUGAGA	1410	TCTCACC C TCATCCTT	2134
1557	AAAGGAUG CUGAUGAG X CGAA IGGGUGAG	1411	CTCACCC T CATCCTT	2135
1559	UAAAAGGA CUGAUGAG X CGAA IAGGGGUG	1412	CACCCCTC A TCCTTTTA	2136
1562	AAGUAAA CUGAUGAG X CGAA IAUGAGGG	1413	CCCTCATC C TTTACTT	2137
1563	AAAGUAAA CUGAUGAG X CGAA IGAUGAGG	1414	CCTCATCC T TTTACTTT	2138
1569	GGGCAAAA CUGAUGAG X CGAA IUAAAAGG	1415	CCTTTTAC T TTTGCCC	2139
1576	GUGGAAGG CUGAUGAG X CGAA ICAAAAAG	1416	CTTTTGC C CCTCCAC	2140
1577	AGUGGAAG CUGAUGAG X CGAA ICAAAAAA	1417	TTTTTGCC C CTTCCACT	2141
1578	AAGUGGAA CUGAUGAG X CGAA IGGCAAAA	1418	TTTTGCCC C TTCCACTT	2142
1579	AAAGUGGA CUGAUGAG X CGAA IGGGCAAA	1419	TTTGCCCC T TCCACTTT	2143
1582	CUCAAGU CUGAUGAG X CGAA IAAGGGGC	1420	GCCCCTTC C ACTTTGAG	2144
1583	ACUCAAG CUGAUGAG X CGAA IGAAGGGG	1421	CCCCTTCC A CTTTGAGT	2145
1585	GUACUCAA CUGAUGAG X CGAA IUGGAAGG	1422	CCTTCCAC T TTGAGTAC	2146
1594	GUGGAUUU CUGAUGAG X CGAA IUACUCAA	1423	TTGAGTAC C AAATCCAC	2147
1595	UGUGGAUU CUGAUGAG X CGAA IGUACUCA	1424	TGAGTACC A AATCCACA	2148
1600	UGGCUUGU CUGAUGAG X CGAA IAUUUGGU	1425	ACCAAATC C ACAAGCCA	2149
1601	AUGGCUUG CUGAUGAG X CGAA IGAUUUGG	1426	CCAAATCC A CAAGCCAT	2150
1603	AAAUGGCU CUGAUGAG X CGAA IUGGAUUU	1427	AAATCCAC A AGCCATTT	2151
1607	CAAAAAAU CUGAUGAG X CGAA ICUUGUGG	1428	CCACAAGC C ATTTTGTG	2152

Table 4

1608	UCAAAAAA CUGAUGAG X CGAA IGCUGUG	1429	CACAAGCC A TTTTTTGA	2153
1636	GCCAGCAU CUGAUGAG X CGAA IUACUCUC	1430	GAGAGTAC C ATGCTGGC	2154
1637	CGCCAGCA CUGAUGAG X CGAA IGUACUCU	1431	AGAGTACC A TGCTGGCG	2155
1641	GCGCCGCC CUGAUGAG X CGAA ICAUGGUA	1432	TACCATGC T GCGGCGC	2156
1650	CUUCCCUC CUGAUGAG X CGAA ICGCCGCC	1433	GGCGGCGC A GAGGGAAG	2157
1663	CGGGUGUA CUGAUGAG X CGAA ICCCUUC	1434	GAAGGGGC C TACACCCG	2158
1664	ACGGGUGU CUGAUGAG X CGAA IGCCCUU	1435	AAGGGGCC T ACACCCGT	2159
1667	AAGACGGG CUGAUGAG X CGAA IUAGGCC	1436	GGGCTTAC A CCCGTCTT	2160
1669	CCAAGACG CUGAUGAG X CGAA IUGUAGG	1437	GCCTACAC C CGTCTTGG	2161
1670	CCAAGAC CUGAUGAG X CGAA IGUGUAGG	1438	CCTACACC C GTCTTGGG	2162
1674	GAGCCCCA CUGAUGAG X CGAA IACGGGUG	1439	CACCCGTC T TGGGGCTC	2163
1681	GUGGGGCG CUGAUGAG X CGAA ICCCAAG	1440	CTTGGGGC T CGCCCCAC	2164
1685	CUGGGUGG CUGAUGAG X CGAA ICGAGCCC	1441	GGGCTCGC C CCACCCAG	2165
1686	CCUGGGUG CUGAUGAG X CGAA ICGAGCC	1442	GGCTCGCC C CACCCAGG	2166
1687	CCCUGGGU CUGAUGAG X CGAA IGGCGAGC	1443	GCTCGCCC C ACCCAGGG	2167
1688	GCCCUGGG CUGAUGAG X CGAA IGGCGAG	1444	CTCGCCCC A CCCAGGGC	2168
1690	GAGCCCUG CUGAUGAG X CGAA IUGGGGCG	1445	CGCCCCAC C CAGGGCTC	2169
1691	GGAGCCCU CUGAUGAG X CGAA IGUGGGGC	1446	GCCCCACC C AGGGCTCC	2170
1692	GGGAGCCC CUGAUGAG X CGAA IGUGGGG	1447	CCCCACC A GGGCTCCC	2171
1697	CAGGAGGG CUGAUGAG X CGAA ICCUGGG	1448	CCCAGGGC T CCCTCTG	2172
1699	UCCAGGAG CUGAUGAG X CGAA IAGCCUG	1449	CAGGGCTC C CTCCTGGA	2173
1700	CUCCAGGA CUGAUGAG X CGAA IGAGCCCU	1450	AGGGCTCC C TCCTGGAG	2174
1701	GCUCCAGG CUGAUGAG X CGAA IGGAGCCC	1451	GGGCTCCC T CCTGGAGC	2175
1703	AUGCUGCA CUGAUGAG X CGAA IAGGGAGC	1452	GCTCCCTC C TGGAGCAT	2176
1704	GAUGCUGC CUGAUGAG X CGAA IAGGGAG	1453	CTCCCTCC T GGAGCATC	2177
1710	GCCUGGGA CUGAUGAG X CGAA ICUCCAGG	1454	CCTGGAGC A TCCCAGGC	2178
1713	CCCGCCUG CUGAUGAG X CGAA IAUUCUC	1455	GGAGCATC C CAGGCGGG	2179
1714	GCCCGCCU CUGAUGAG X CGAA IGAUCUC	1456	GAGCATCC C AGGCGGGC	2180
1715	CGCCCGCC CUGAUGAG X CGAA IGAUGCU	1457	AGCATCCC A GCGGGCG	2181
1726	GUCUGGCG CUGAUGAG X CGAA ICCGCCCC	1458	CGGGCGGC A CGCCAGAC	2182
1730	GGCUGUCU CUGAUGAG X CGAA ICGUGCCG	1459	CGGCACGC C AGACAGCC	2183
1731	GGGUGUC CUGAUGAG X CGAA ICGUGCC	1460	GGCACGCC A GACAGCCC	2184
1735	GGGGGGGC CUGAUGAG X CGAA IUCUGGCG	1461	CGCCAGAC A GCCCCCCC	2185
1738	AAGGGGGG CUGAUGAG X CGAA ICUGUCUG	1462	CAGACAGC C CCCCCCTT	2186
1739	CAAGGGGG CUGAUGAG X CGAA ICGUGUCU	1463	AGACAGCC C CCCCCCTG	2187
1740	UCAAGGGG CUGAUGAG X CGAA IGGCUGUC	1464	GACAGCCC C CCCCTTGA	2188
1741	UUCAAGGG CUGAUGAG X CGAA IGGCUGU	1465	ACAGCCCC C CCCTTGAA	2189
1742	AUJCAAGG CUGAUGAG X CGAA IGGGCUG	1466	CAGCCCCC C CCTTGAAT	2190
1743	GAUJCAAG CUGAUGAG X CGAA IGGGGCU	1467	AGCCCCC C CTTGAATC	2191
1744	AGAUJCAA CUGAUGAG X CGAA IGGGGGC	1468	GCCCCCC C TTGAATCT	2192
1745	CAGAUJCA CUGAUGAG X CGAA IGGGGGG	1469	CCCCCCC T TGAATCTG	2193
1752	CUCCUGC CUGAUGAG X CGAA IAUJCAAG	1470	CTTGAATC T GCAGGGAG	2194
1755	UUGCUC CUGAUGAG X CGAA ICAGAUUC	1471	GAATCTGC A GGGAGCAA	2195
1762	UGGAGAGU CUGAUGAG X CGAA ICUCCUG	1472	CAGGGAGC A ACTCTCCA	2196
1765	GAGUGGAG CUGAUGAG X CGAA IUUGCUC	1473	GGAGCAAC T CTCCACTC	2197
1767	UGGAGUGG CUGAUGAG X CGAA IAGUUGCU	1474	AGCAACTC T CCACTCCA	2198
1769	UAUGGAGU CUGAUGAG X CGAA IAGAGUUG	1475	CAACTCTC C ACTCCATA	2199

Table 4

1770	AUAUGGAG CUGAUGAG X CGAA IGAGAGUU	1476	AACTCTCC A CTCCATAT	2200
1772	AAAUAUGG CUGAUGAG X CGAA IUGGAGAG	1477	CTCTCCAC T CCATATTT	2201
1774	AUAAAUAU CUGAUGAG X CGAA IAGUGGAG	1478	CTCCACTC C ATATTTAT	2202
1775	AAUAAAUA CUGAUGAG X CGAA IGAGUGGA	1479	TCCACTCC A TATTTATT	2203
1789	GAAAAAAU CUGAUGAG X CGAA IUUUAAAU	1480	ATTTAAAC A ATTTTTC	2204
1798	GCCUUUGG CUGAUGAG X CGAA IAAAAAU	1481	ATTTTTC C CCAAAGGC	2205
1799	UGCCUUUG CUGAUGAG X CGAA IGAAAAA	1482	TTTTTCC C CAAAGGCA	2206
1800	AUGCCUUU CUGAUGAG X CGAA IGGAAAA	1483	TTTTTCCC C AAAGGCAT	2207
1801	GAUGCCUU CUGAUGAG X CGAA IGGAAAA	1484	TTTTCCCC A AAGGCATC	2208
1807	ACUAUGGA CUGAUGAG X CGAA ICCUUUGG	1485	CCAAAGGC A TCCATAGT	2209
1810	UGCACUUA CUGAUGAG X CGAA IAUGCCUU	1486	AAGGCATC C ATAGTGCA	2210
1811	GUGCACUA CUGAUGAG X CGAA IGAUGCCU	1487	AGGCATCC A TAGTGCAC	2211
1818	AAUGCUG CUGAUGAG X CGAA ICACUAUG	1488	CATAGTGC A CTAGCATT	2212
1820	AAAAUGCU CUGAUGAG X CGAA IUGCACUA	1489	TAGTGCAC T AGCATTTT	2213
1824	CAAGAAAA CUGAUGAG X CGAA ICUAGUGC	1490	GCACTAGC A TTTTCTTG	2214
1830	UUGGUUCA CUGAUGAG X CGAA IAAAAUGC	1491	GCATTTTC T TGAACCAA	2215
1836	ACAUAUUA CUGAUGAG X CGAA IUUCAAGA	1492	TCTTGAAC C AATAATGT	2216
1837	UACAUUAU CUGAUGAG X CGAA IGUUCAAG	1493	CTTGAACC A ATAATGTA	2217
1864	UGCAAGGC CUGAUGAG X CGAA IACAUCAA	1494	TTGATGTC A GCCTTGCA	2218
1867	UGAUGCAA CUGAUGAG X CGAA ICUGACAU	1495	ATGTCAGC C TTGCATCA	2219
1868	UUGAUGCA CUGAUGAG X CGAA IGCUGACA	1496	TGTCAGCC T TGCATCAA	2220
1872	GCCCUUGA CUGAUGAG X CGAA ICAAGGCU	1497	AGCCTTGC A TCAAGGGC	2221
1875	AAAGCCCU CUGAUGAG X CGAA IAUGCAAG	1498	CTTGCATC A AGGGCTTT	2222
1881	UUUGAUAA CUGAUGAG X CGAA ICCCUUGA	1499	TCAAGGGC T TTATCAAA	2223
1887	GUACUUUU CUGAUGAG X CGAA IAUAAAGC	1500	GCTTTATC A AAAAGTAC	2224
1896	UUUAUUUA CUGAUGAG X CGAA IUACUUUU	1501	AAAAGTAC A ATAATAAA	2225
1907	CUACCUGA CUGAUGAG X CGAA IAUUUUAU	1502	AATAAATC C TCAGGTAG	2226
1908	ACUACCUG CUGAUGAG X CGAA IGAUUUAU	1503	ATAAATCC T CAGGTAGT	2227
1910	GUACUACC CUGAUGAG X CGAA IAGGAUUU	1504	AAATCCTC A GGTAGTAC	2228
1919	CCAUUCCC CUGAUGAG X CGAA IUACUACC	1505	GGTAGTAC T GGAATGG	2229
1933	CAUGGCAA CUGAUGAG X CGAA ICCUCCA	1506	TGGAAGGC T TTGCCATG	2230
1938	AGGCCCAU CUGAUGAG X CGAA ICAAAGCC	1507	GGCTTGC C ATGGGCCT	2231
1939	CAGGCCCA CUGAUGAG X CGAA IGCAAAGC	1508	GCTTTGCC A TGGGCCTG	2232
1945	ACGAGCA CUGAUGAG X CGAA ICCCAUGG	1509	CCATGGGC C TGCTGCGT	2233
1946	GACGAGC CUGAUGAG X CGAA IGCCCAUG	1510	CATGGGCC T GCTGCGTC	2234
1949	UCUGACGC CUGAUGAG X CGAA ICAGGCC	1511	GGGCCTGC T GCGTCAGA	2235
1955	UACUGGUC CUGAUGAG X CGAA IACGCAGC	1512	GCTGCGTC A GACCAGTA	2236
1959	CCAGUACU CUGAUGAG X CGAA IUCUGACG	1513	CGTCAGAC C AGTACTGG	2237
1960	CCCAGUAC CUGAUGAG X CGAA IGUCUGAC	1514	GTCAGACC A GTACTGGG	2238
1965	UCCUUCCC CUGAUGAG X CGAA IUACUGGU	1515	ACCAGTAC T GGAAGGA	2239
1988	AUAACAAC CUGAUGAG X CGAA ICUUACAA	1516	TTGTAAGC A GTTGTAT	2240
2032	UAUAGCAU CUGAUGAG X CGAA IUUCUAUC	1517	GATAGAAC A ATGCTATA	2241
2037	UAUAUUUA CUGAUGAG X CGAA ICAUUGUU	1518	AACAATGC T ATAATATA	2242
2054	UACCCACG CUGAUGAG X CGAA IUUCAUUA	1519	TAATGAAC A CGTGGGTA	2243
2076	UCACAUCA CUGAUGAG X CGAA IUUUCUUA	1520	TAAGAAAC A TGATGTGA	2244
2091	CGGGACAA CUGAUGAG X CGAA IUAUCUC	1521	GAGATTAC T TTGTCCCG	2245
2097	AAUAAGCG CUGAUGAG X CGAA IACAAAGU	1522	ACTTTGTC C CGTTATT	2246

Table 4

2098	GAAUAAGC	CUGAUGAG	X	CGAA	IGACAAAG	1523	CTTTGTCC	C	GCTTATTC	2247
2101	GCAGAAUA	CUGAUGAG	X	CGAA	ICGGGACA	1524	TGTCCCGC	T	TATTCTGC	2248
2107	CAGGGAGC	CUGAUGAG	X	CGAA	IAAUAAGC	1525	GCTTATTC	T	GCTCCCTG	2249
2110	UAACAGGG	CUGAUGAG	X	CGAA	ICAGAAUA	1526	TATTCTGC	T	CCCTGTTA	2250
2112	GAUACAG	CUGAUGAG	X	CGAA	IAGCAGAA	1527	TTCTGCTC	C	CTGTTATC	2251
2113	AGAUAACA	CUGAUGAG	X	CGAA	IGAGCAGA	1528	TCTGCTCC	C	TGTTATCT	2252
2114	CAGAUAAC	CUGAUGAG	X	CGAA	IGGAGCAG	1529	CTGCTCCC	T	GTTATCTG	2253
2121	GAUCUAGC	CUGAUGAG	X	CGAA	IAUAACAG	1530	CTGTTATC	T	GCTAGATC	2254
2124	CUAGAUUC	CUGAUGAG	X	CGAA	ICAGAUAA	1531	TTATCTGC	T	AGATCTAG	2255
2130	UGAGAACU	CUGAUGAG	X	CGAA	IAUCUAGC	1532	GCTAGATC	T	AGTTCTCA	2256
2136	AGUGAUUG	CUGAUGAG	X	CGAA	IAACUAGA	1533	TCTAGTTC	T	CAATCACT	2257
2138	GCAGUGAU	CUGAUGAG	X	CGAA	IAGAACUA	1534	TAGTTCTC	A	ATCACTGC	2258
2142	GGGAGCAG	CUGAUGAG	X	CGAA	IAUUGAGA	1535	TCTCAATC	A	CTGCTCCC	2259
2144	GGGGAGC	CUGAUGAG	X	CGAA	IUGAUUGA	1536	TCAATCAC	T	GCTCCCCC	2260
2147	CACGGGGG	CUGAUGAG	X	CGAA	ICAGUGAU	1537	ATCACTGC	T	CCCCCGTG	2261
2149	CACACGGG	CUGAUGAG	X	CGAA	IAGCAGUG	1538	CACTGCTC	C	CCCGTGTG	2262
2150	ACACACGG	CUGAUGAG	X	CGAA	IGAGCAGU	1539	ACTGCTCC	C	CGTGTGT	2263
2151	UACACACG	CUGAUGAG	X	CGAA	IGGAGCAG	1540	CTGCTCCC	C	CGTGTGTA	2264
2152	AUACACAC	CUGAUGAG	X	CGAA	IGGGAGCA	1541	TGCTCCCC	C	GTGTGTAT	2265
2169	ACCUUACA	CUGAUGAG	X	CGAA	ICAUUCUA	1542	TAGAATGC	A	TGTAAGGT	2266
2179	ACACAAGA	CUGAUGAG	X	CGAA	IACCUUAC	1543	GTAAGGTC	T	TCTGTGT	2267
2182	AGGACACA	CUGAUGAG	X	CGAA	IAAGACCU	1544	AGGTCTTC	T	TGTGTCCT	2268
2189	UUUCAUCA	CUGAUGAG	X	CGAA	IACACAAG	1545	CTTGTGTC	C	TGATGAAA	2269
2190	UUUUAUC	CUGAUGAG	X	CGAA	IGACACAA	1546	TTGTGTCC	T	GATGAAA	2270
2207	UCAUUUCA	CUGAUGAG	X	CGAA	ICACAUAU	1547	ATATGTGC	T	TGAAATGA	2271
2221	GAGAUCAA	CUGAUGAG	X	CGAA	IUUUCUA	1548	TGAGAAAC	T	TTGATCTC	2272
2228	GUAAGCAG	CUGAUGAG	X	CGAA	IAUCAAAG	1549	CTTTGATC	T	CTGCTTAC	2273
2230	UAGUAAGC	CUGAUGAG	X	CGAA	IAGAUCAA	1550	TTGATCTC	T	GCTTACTA	2274
2233	CAUUAGUA	CUGAUGAG	X	CGAA	ICAGAGAU	1551	ATCTCTGC	T	TACTAATG	2275
2237	GGCACAUU	CUGAUGAG	X	CGAA	IUAAGCAG	1552	CTGCTTAC	T	AATGTGCC	2276
2245	GGACAUGG	CUGAUGAG	X	CGAA	ICACAUUA	1553	TAATGTGC	C	CCATGTCC	2277
2246	UGGACAUG	CUGAUGAG	X	CGAA	IGCACAUU	1554	AATGTGCC	C	CATGTCCA	2278
2247	UUGGACAU	CUGAUGAG	X	CGAA	IGGCACAU	1555	ATGTGCCC	C	ATGTCCAA	2279
2248	CUUGGACA	CUGAUGAG	X	CGAA	IGGGCACA	1556	TGTGCCCC	A	TGTCCAAG	2280
2253	UUGGACUU	CUGAUGAG	X	CGAA	IACAUGGG	1557	CCCATGTC	C	AAGTCCAA	2281
2254	GUUGGACU	CUGAUGAG	X	CGAA	IGACAUGG	1558	CCATGTCC	A	AGTCCAAC	2282
2259	GGCAGGUU	CUGAUGAG	X	CGAA	IACUUGGA	1559	TCCAAGTC	C	AACCTGCC	2283
2260	AGGCAGGU	CUGAUGAG	X	CGAA	IGACUUGG	1560	CCAAGTCC	A	ACCTGCCT	2284
2263	CACAGGCA	CUGAUGAG	X	CGAA	IUUGGACU	1561	AGTCCAAC	C	TGCCTGTG	2285
2264	GCACAGGC	CUGAUGAG	X	CGAA	IGUUGGAC	1562	GTCCAACC	T	GCCTGTGC	2286
2267	CAUGCACA	CUGAUGAG	X	CGAA	ICAGGUUG	1563	CAACCTGC	C	TGTGCATG	2287
2268	UCAUGCAC	CUGAUGAG	X	CGAA	IGCAGGUU	1564	AACCTGCC	T	GTGCATGA	2288
2273	UCAGGUCA	CUGAUGAG	X	CGAA	ICACAGGC	1565	GCCTGTGC	A	TGACCTGA	2289
2278	AAUGAUCA	CUGAUGAG	X	CGAA	IUCAUGCA	1566	TGCATGAC	C	TGATCATT	2290
2279	UAAUGAUC	CUGAUGAG	X	CGAA	IGUCAUGC	1567	GCATGACC	T	GATCATT	2291
2284	CCAUGUAA	CUGAUGAG	X	CGAA	IAUCAGGU	1568	ACCTGATC	A	TTACATGG	2292
2289	CACAGCCA	CUGAUGAG	X	CGAA	IUAAUGAU	1569	ATCATTAC	A	TGGCTGTG	2293

Table 4

2294	GGAACCAC CUGAUGAG X CGAA ICCAUGUA	1570	TACATGGC T GTGGTTCC	2294
2302	CAGGCUUA CUGAUGAG X CGAA IAACCACA	1571	TGTGGTTC C TAAGCCTG	2295
2303	ACAGGCUU CUGAUGAG X CGAA IGAACCAC	1572	GTGGTTCC T AAGCCTGT	2296
2308	CAGCAACA CUGAUGAG X CGAA ICUUAGGA	1573	TCCTAAGC C TGTGCTG	2297
2309	UCAGCAAC CUGAUGAG X CGAA IGCUUAGG	1574	CCTAAGCC T GTTGCTGA	2298
2315	AUGACUUC CUGAUGAG X CGAA ICAACAGG	1575	CCTGTTGC T GAAGTCAT	2299
2322	AGCGACAA CUGAUGAG X CGAA IACUUCAG	1576	CTGAAGTC A TTGTCGCT	2300
2330	UAUUGCUG CUGAUGAG X CGAA ICGACAAU	1577	ATTGTCGC T CAGCAATA	2301
2332	CCUAUUGC CUGAUGAG X CGAA IAGCGACA	1578	TGTCGCTC A GCAATAGG	2302
2335	CACCCUUA CUGAUGAG X CGAA ICUGAGCG	1579	CGCTCAGC A ATAGGCTG	2303
2345	UGGAAAAC CUGAUGAG X CGAA ICACCCUA	1580	TAGGGTGC A GTTTTCCA	2304
2352	CUAUUCCU CUGAUGAG X CGAA IAAAACUG	1581	CAGTTTTC C AGGAATAG	2305
2353	CCUAUUCC CUGAUGAG X CGAA IGAAAACU	1582	AGTTTTC A GGAATAGG	2306
2363	UAGGCAAA CUGAUGAG X CGAA ICCUAUUC	1583	GAATAGGC A TTTGCCTA	2307
2369	AGGAAUUA CUGAUGAG X CGAA ICAAAUGC	1584	GCATTGTC C TAATTCCT	2308
2370	CAGGAAUU CUGAUGAG X CGAA IGCAAUG	1585	CATTGCCC T AATTCCTG	2309
2376	UCAUGCCA CUGAUGAG X CGAA IAAUUAGG	1586	CCTAATTC C TGGCATGA	2310
2377	GUCAUGCC CUGAUGAG X CGAA IGAAUUAG	1587	CTAATTC T GGCATGAC	2311
2381	GAGUGUCA CUGAUGAG X CGAA ICCAGGAA	1588	TTCTGGC A TGACACTC	2312
2386	CACUAGAG CUGAUGAG X CGAA IUCAUGCC	1589	GGCATGAC A CTCTAGTG	2313
2388	GUCACUAG CUGAUGAG X CGAA IUGUCAUG	1590	CATGACAC T CTAGTGAC	2314
2390	AAGUCACU CUGAUGAG X CGAA IAGUGUCA	1591	TGACACTC T AGTGACTT	2315
2397	CACCAGGA CUGAUGAG X CGAA IUCACUAG	1592	CTAGTGAC T TCCTGGTG	2316
2400	CCUCACCA CUGAUGAG X CGAA IAAGUCAC	1593	GTGACTTC C TGGTGAGG	2317
2401	GCCUCACC CUGAUGAG X CGAA IGAAGUCA	1594	TGACTTCC T GGTGAGGC	2318
2410	ACAGGCUG CUGAUGAG X CGAA ICCUCACC	1595	GGTGAGGC C CAGCCTGT	2319
2411	GACAGGCU CUGAUGAG X CGAA IGCCUCAC	1596	GTGAGGCC C AGCCTGTC	2320
2412	GGACAGGC CUGAUGAG X CGAA IGGCCUCA	1597	TGAGGCC A GCCTGTCC	2321
2415	CCAGGACA CUGAUGAG X CGAA ICUGGGCC	1598	GGCCAGC C TGCTCTGG	2322
2416	ACCAGGAC CUGAUGAG X CGAA IGCUGGGC	1599	GCCCAGCC T GTCCTGGT	2323
2420	CUGUACCA CUGAUGAG X CGAA IACAGGCU	1600	AGCCTGTC C TGGTACAG	2324
2421	GCUGUACC CUGAUGAG X CGAA IGACAGGC	1601	GCCTGTCC T GGTACAGC	2325
2427	GACCCUGC CUGAUGAG X CGAA IUACCAGG	1602	CCTGGTAC A GCAGGCTC	2326
2430	CAAGACCC CUGAUGAG X CGAA ICUGUACC	1603	GGTACAGC A GGTCTTG	2327
2436	UUACAGCA CUGAUGAG X CGAA IACCCUGC	1604	GCAGGGTC T TGCTGTAA	2328
2440	UGAGUUAC CUGAUGAG X CGAA ICAAGACC	1605	GGTCTTGC T GTAACCTA	2329
2446	AAUGUCUG CUGAUGAG X CGAA IUUACAGC	1606	GCTGTAA C CAGACATT	2330
2448	GGAAUGUC CUGAUGAG X CGAA IAGUUACA	1607	TGTAACCT A GACATTCC	2331
2452	CCUUGGAA CUGAUGAG X CGAA IUCUGAGU	1608	ACTCAGAC A TTCCAAGG	2332
2456	AUACCCUU CUGAUGAG X CGAA IAAUGUCU	1609	AGACATTC C AAGGGTAT	2333
2457	CAUACCCU CUGAUGAG X CGAA IGAAUGUC	1610	GACATTCC A AGGGTATG	2334
2472	GUGAAUAU CUGAUGAG X CGAA ICUUCCCA	1611	TGGGAAGC C ATATTCAC	2335
2473	UGUGAAUA CUGAUGAG X CGAA IGCUUCCC	1612	GGGAAGCC A TATTCACA	2336
2479	GUGAGGUG CUGAUGAG X CGAA IAAUAUGG	1613	CCATATTC A CACCTCAC	2337
2481	GCGUGAGG CUGAUGAG X CGAA IUGAAUAU	1614	ATATTCAC A CCTCACGC	2338
2483	GAGCGUGA CUGAUGAG X CGAA IUGUGAAU	1615	ATTACAC C TCACGCTC	2339
2484	AGAGCGUG CUGAUGAG X CGAA IGUGUGAA	1616	TTACACCC T CACGCTCT	2340

Table 4

2486	CCAGAGCG CUGAUGAG X CGAA IAGGUGUG	1617	CACACCTC A CGCTCTGG	2341
2490	AUGUCCAG CUGAUGAG X CGAA ICGUGAGG	1618	CCTCACGC T CTGGACAT	2342
2492	UCAUGUCC CUGAUGAG X CGAA IAGCGUGA	1619	TCACGCTC T GGACATGA	2343
2497	CUAAAUCA CUGAUGAG X CGAA IUCCAGAG	1620	CTCTGGAC A TGATTTAG	2344
2512	GGUGUCCC CUGAUGAG X CGAA ICUUCCCU	1621	AGGGAAGC A GGGACACC	2345
2518	GCGGGGGG CUGAUGAG X CGAA IUCCUGC	1622	GCAGGGAC A CCCCCCGC	2346
2520	GGGCGGGG CUGAUGAG X CGAA IUGUCCCU	1623	AGGGACAC C CCCCCCCC	2347
2521	GGGGCGGG CUGAUGAG X CGAA IGUGUCCC	1624	GGGACACC C CCGCCCCC	2348
2522	GGGGGCGG CUGAUGAG X CGAA IGGUGUCC	1625	GGACACCC C CCGCCCCC	2349
2523	GGGGGGCG CUGAUGAG X CGAA IGGGUGUC	1626	GACACCCC C CGCCCCC	2350
2524	UGGGGGGC CUGAUGAG X CGAA IGGGGUGU	1627	ACACCCCC C GCCCCCA	2351
2527	AGGUGGGG CUGAUGAG X CGAA ICGGGGGG	1628	CCCCCGC C CCCCACCT	2352
2528	AAGGUGGG CUGAUGAG X CGAA ICGGGGGG	1629	CCCCCGCC C CCCACCTT	2353
2529	AAAGGUGG CUGAUGAG X CGAA IGGCGGGG	1630	CCCCGCCC C CCACCTTT	2354
2530	CAAAGGUG CUGAUGAG X CGAA IGGCGGGG	1631	CCCGCCCC C CACCTTTG	2355
2531	CCAAAGGU CUGAUGAG X CGAA IGGGCGGG	1632	CCGCCCCC C ACCTTTGG	2356
2532	CCCAAAGG CUGAUGAG X CGAA IGGGGGCG	1633	CGCCCCC A CCTTTGGG	2357
2534	AUCCCAA CUGAUGAG X CGAA IUGGGGGG	1634	CCCCCAC C TTGGGAT	2358
2535	GAUCCCAA CUGAUGAG X CGAA IGUGGGGG	1635	CCCCCACC T TTGGGATC	2359
2544	GCGGAGGC CUGAUGAG X CGAA IAUCCCAA	1636	TTGGGATC A GCCTCCGC	2360
2547	AUGGCGGA CUGAUGAG X CGAA ICUGAUCC	1637	GGATCAGC C TCCGCCAT	2361
2548	AAUGGCGG CUGAUGAG X CGAA IGCUGAUC	1638	GATCAGCC T CCGCCATT	2362
2550	GGAAUGGC CUGAUGAG X CGAA IAGGCUGA	1639	TCAGCCTC C GCCATTCC	2363
2553	CUUGGAAU CUGAUGAG X CGAA ICGGAGGC	1640	GCCTCCGC C ATTCCAAG	2364
2554	ACUUGGAA CUGAUGAG X CGAA ICGGAGG	1641	CCTCCGCC A TTCCAAGT	2365
2558	GUCGACUU CUGAUGAG X CGAA IAAUGGCG	1642	CGCCATTC C AAGTCGAC	2366
2559	UGUCGACU CUGAUGAG X CGAA IGAUUGGC	1643	GCCATTCC A AGTCGACA	2367
2567	AAGAAGAG CUGAUGAG X CGAA IUCGACUU	1644	AAGTCGAC A CTCTTCTT	2368
2569	UCAAGAAG CUGAUGAG X CGAA IUGUCGAC	1645	GTCGACAC T CTCTTGA	2369
2571	GCUCAAGA CUGAUGAG X CGAA IAGUGUCG	1646	CGACACTC T TCTTGAGC	2370
2574	UCUGCUCA CUGAUGAG X CGAA IAAGAGUG	1647	CACTCTTC T TGAGCAGA	2371
2580	UCACGGUC CUGAUGAG X CGAA ICUCAAGA	1648	TCTTGAGC A GACCGTGA	2372
2584	CAAUAC CUGAUGAG X CGAA IUCUGCUC	1649	GAGCAGAC C GTGATTG	2373
2603	CCAGCAGG CUGAUGAG X CGAA ICCUCUCU	1650	AGAGAGGC A CCGCTGG	2374
2605	UUCCAGCA CUGAUGAG X CGAA IUGCCUCU	1651	AGAGGCAC C TGCTGGAA	2375
2606	UUUCCAGC CUGAUGAG X CGAA IGUGCCUC	1652	GAGGCACC T GCTGAAA	2376
2609	UGGUUCC CUGAUGAG X CGAA ICAGGUGC	1653	GCACCTGC T GGAAACCA	2377
2616	AGAAGUGU CUGAUGAG X CGAA IUUCCAG	1654	CTGGAAAC C ACACTTCT	2378
2617	AAGAAGUG CUGAUGAG X CGAA IGUUCCA	1655	TGGAAACC A CACTTCTT	2379
2619	UCAAGAAG CUGAUGAG X CGAA IUGGUUUC	1656	GAAACCAC A CTCTTGA	2380
2621	UUUCAAGA CUGAUGAG X CGAA IUGUGGUU	1657	AACCACAC T TCTGAAA	2381
2624	CUGUUUCA CUGAUGAG X CGAA IAAGUGUG	1658	CACACTTC T TGAAACAG	2382
2631	ACCCAGGC CUGAUGAG X CGAA IUUUAAG	1659	CTTGAAAC A GCCTGGGT	2383
2634	GUCACCCA CUGAUGAG X CGAA ICUGUUUC	1660	GAAACAGC C TGGTGAC	2384
2635	CGUCACCC CUGAUGAG X CGAA IGCUGUUU	1661	AAACAGCC T GGGTGACG	2385
2647	UGCCUAAA CUGAUGAG X CGAA IACCGUCA	1662	TGACGGTC C TTTAGGCA	2386
2648	CUGCCUAA CUGAUGAG X CGAA IGACCGUC	1663	GACGGTCC T TTAGGCAG	2387

Table 4

2655	CGGCAGGC CUGAUGAG X CGAA ICCUAAAG	1664	CTTTAGGC A GCCTGCCG	2388
2658	CGGCGGCA CUGAUGAG X CGAA ICUGCCUA	1665	TAGGCAGC C TGCCGCCG	2389
2659	ACGGCGGC CUGAUGAG X CGAA IGCUGCCU	1666	AGGCAGCC T GCCGCCGT	2390
2662	GAGACGGC CUGAUGAG X CGAA ICAGGCUG	1667	CAGCCTGC C GCCGTCTC	2391
2665	ACAGAGAC CUGAUGAG X CGAA ICGGCAGG	1668	CCTGCCGC C GTCTCTGT	2392
2669	CGGGACAG CUGAUGAG X CGAA IACGGCGG	1669	CCGCCGTC T CTGTCCCG	2393
2671	ACCGGGAC CUGAUGAG X CGAA IAGACGGC	1670	GCCGTCTC T GTCCCGT	2394
2675	GUGAACCG CUGAUGAG X CGAA IACAGAGA	1671	TCTCTGTC C CGGTTTAC	2395
2676	GGUGAACC CUGAUGAG X CGAA IGACAGAG	1672	CTCTGTCC C GGTTCACC	2396
2682	CGGCAAGG CUGAUGAG X CGAA IAACCGGG	1673	CCCGGTTT A CCTTGCCG	2397
2684	CUCGGCAA CUGAUGAG X CGAA IUGAACCG	1674	CGGTTTAC C TTGCCGAG	2398
2685	UCUCGGCA CUGAUGAG X CGAA IGUGAACC	1675	GGTTTACC T TGCCGAGA	2399
2689	CCUCUCUC CUGAUGAG X CGAA ICAAGGUG	1676	CACCTTGC C GAGAGAGG	2400
2704	GGUGGGGC CUGAUGAG X CGAA IACGCGCC	1677	GGCGCGTC T GCCCACC	2401
2707	GAGGGUGG CUGAUGAG X CGAA ICAGACGC	1678	GCGTCTGC C CCACCCTC	2402
2708	UGAGGGUG CUGAUGAG X CGAA IGCAGACG	1679	CGTCTGCC C CACCCTCA	2403
2709	UUGAGGGU CUGAUGAG X CGAA IGGCAGAC	1680	GTCTGCCC C ACCCTCAA	2404
2710	UUUGAGGG CUGAUGAG X CGAA IGGGCAGA	1681	TCTGCCCC A CCCTCAA	2405
2712	GGUUUGAG CUGAUGAG X CGAA IUGGGGCA	1682	TGCCCCAC C CTCAAACC	2406
2713	GGGUUGA CUGAUGAG X CGAA IGUGGGGC	1683	GCCCCACC C TCAAACCC	2407
2714	AGGGUUUG CUGAUGAG X CGAA IGGUGGGG	1684	CCCCACC T CAAACCCT	2408
2716	ACAGGGUU CUGAUGAG X CGAA IAGGGUGG	1685	CCACCCTC A AACCTGT	2409
2720	CCCCACAG CUGAUGAG X CGAA IUUUGAGG	1686	CCTCAAAC C CTGTGGGG	2410
2721	GCCCCACA CUGAUGAG X CGAA IGUUUGAG	1687	CTCAAACC C TGTGGGGC	2411
2722	GGCCCCAC CUGAUGAG X CGAA IGGUUUGA	1688	TCAAACCC T GTGGGGCC	2412
2730	CACCAUCA CUGAUGAG X CGAA ICCCCACA	1689	TGTGGGGC C TGATGGTG	2413
2731	GCACCAUC CUGAUGAG X CGAA IGCCCCAC	1690	GTGGGGCC T GATGGTGC	2414
2740	GAGUCGUG CUGAUGAG X CGAA ICACCAUC	1691	GATGGTGC T CACGACTC	2415
2742	AAGAGUCG CUGAUGAG X CGAA IAGCACCA	1692	TGGTGCTC A CGACTCTT	2416
2747	GCAGGAAG CUGAUGAG X CGAA IUCGUGAG	1693	CTCACGAC T CTTCTTGC	2417
2749	UUGCAGGA CUGAUGAG X CGAA IAGUCGUG	1694	CACGACTC T TCCTGCAA	2418
2752	CCUUUGCA CUGAUGAG X CGAA IAAGAGUC	1695	GACTCTTC C TGCAAAGG	2419
2753	CCCUUUGC CUGAUGAG X CGAA IGAAGAGU	1696	ACTCTTCC T GCAAAGGG	2420
2756	GUUCCCUU CUGAUGAG X CGAA ICAGGAAG	1697	CTTCCTGC A AAGGGAAC	2421
2765	AGGUCUUC CUGAUGAG X CGAA IUUCCCUU	1698	AAGGGAAC T GAAGACCT	2422
2772	AAUGUGGA CUGAUGAG X CGAA IUUCUCAG	1699	CTGAAGAC C TCCACATT	2423
2773	UAAUGUGG CUGAUGAG X CGAA IGUCUUCA	1700	TGAAGACC T CCACATTA	2424
2775	CUUAAUGU CUGAUGAG X CGAA IAGGUCUU	1701	AAGACCTC C ACATTAAAG	2425
2776	ACUUAUUG CUGAUGAG X CGAA IAGGUCUU	1702	AGACCTCC A CATTAAAGT	2426
2778	CCACUUA CUGAUGAG X CGAA IUGGAGGU	1703	ACCTCCAC A TTAAGTGG	2427
2788	UGUUAAAA CUGAUGAG X CGAA ICCACUUA	1704	TAAGTGGC T TTTTAACA	2428
2796	GUUUUUA CUGAUGAG X CGAA IUUAAAAA	1705	TTTTTAAC A TGAAAAAC	2429
2805	AGCUGCCG CUGAUGAG X CGAA IUUUUUA	1706	TGAAAAAC A CGGCAGCT	2430
2810	GCUACAGC CUGAUGAG X CGAA ICCGUGUU	1707	AACACGGC A GCTGTAGC	2431
2813	GGAGCUAC CUGAUGAG X CGAA ICUGCCGU	1708	ACGCAGC T GTAGCTCC	2432
2819	AGCUCGGG CUGAUGAG X CGAA ICUACAGC	1709	GCTGTAGC T CCCGAGCT	2433
2821	GUAGCUCG CUGAUGAG X CGAA IAGCUACA	1710	TGTAGCTC C CGAGCTAC	2434

Table 4

2822	AGUAGCUC CUGAUGAG X CGAA IGAGCUAC	1711	GTAGCTCC C GAGCTACT	2435
2827	AAGAGAGU CUGAUGAG X CGAA ICUCGGGA	1712	TCCCGAGC T ACTCTCTT	2436
2830	GGCAAGAG CUGAUGAG X CGAA IUAGCUCG	1713	CGAGCTAC T CTCTTGCC	2437
2832	CUGGCAAG CUGAUGAG X CGAA IAGUAGCU	1714	AGCTACTC T CTTGCCAG	2438
2834	UGCUGGCA CUGAUGAG X CGAA IAGAGUAG	1715	CTACTCTC T TGCCAGCA	2439
2838	AAAAUGCU CUGAUGAG X CGAA ICAAGAGA	1716	TCTCTTGC C AGCATTTT	2440
2839	GAAAAUGC CUGAUGAG X CGAA IGCAAGAG	1717	CTCTTGCC A GCATTTTC	2441
2842	UGUGAAAA CUGAUGAG X CGAA ICUGGCAA	1718	TTGCCAGC A TTTTCACA	2442
2848	GCAAAAUG CUGAUGAG X CGAA IAAAAUGC	1719	GCATTTTC A CATTTTGC	2443
2850	AGGCAAAA CUGAUGAG X CGAA IUGAAAAU	1720	ATTTTCAC A TTTTGCCT	2444
2857	ACGAGAAA CUGAUGAG X CGAA ICAAAAUG	1721	CATTTTGC C TTTCTCGT	2445
2858	CACGAGAA CUGAUGAG X CGAA IGCAAAAU	1722	ATTTTGCC T TTCTCGTG	2446
2862	CUACCACG CUGAUGAG X CGAA IAAAGGCA	1723	TGCCTTTC T CGTGGTAG	2447
2875	UCUGUACU CUGAUGAG X CGAA ICUUCUAC	1724	GTAGAAGC C AGTACAGA	2448
2876	CUCUGUAC CUGAUGAG X CGAA IGCUCUA	1725	TAGAAGCC A GTACAGAG	2449
2881	AAUUUCUC CUGAUGAG X CGAA IUACUGGC	1726	GCCAGTAC A GAGAAATT	2450
2891	CCCACCAC CUGAUGAG X CGAA IAAUUUCU	1727	AGAAATTC T GTGGTGGG	2451
2903	ACCUCGAA CUGAUGAG X CGAA IUUCCAC	1728	GTGGGAAC A TTCGAGGT	2452
2915	CUGCAGGG CUGAUGAG X CGAA IACACCUC	1729	GAGGTGTC A CCCTGCAG	2453
2917	CUCUGCAG CUGAUGAG X CGAA IUGACACC	1730	GGTGTAC C CTGCAGAG	2454
2918	GCUCUGCA CUGAUGAG X CGAA IGUGACAC	1731	GTGTCACC C TGCAGAGC	2455
2919	AGCUCUGC CUGAUGAG X CGAA IGGUGACA	1732	TGTCACCC T GCAGAGCT	2456
2922	CAUAGCUC CUGAUGAG X CGAA ICAGGGUG	1733	CACCCTGC A GAGCTATG	2457
2927	CUCACCAU CUGAUGAG X CGAA ICUCUGCA	1734	TGCAGAGC T ATGGTGAG	2458
2949	GGCACCUA CUGAUGAG X CGAA ICCUUAUC	1735	GATAAGGC T TAGGTGCC	2459
2957	UACAGCCU CUGAUGAG X CGAA ICACCUAA	1736	TTAGGTGC C AGGCTGTA	2460
2958	UUACAGCC CUGAUGAG X CGAA IGCACCUA	1737	TAGGTGCC A GGCTGTAA	2461
2962	AUGCUUAC CUGAUGAG X CGAA ICCUGGCA	1738	TGCCAGGC T GTAAGCAT	2462
2969	GCUCAGAA CUGAUGAG X CGAA ICUUACAG	1739	CTGTAAGC A TTCTGAGC	2463
2973	GCCAGCUC CUGAUGAG X CGAA IAAUGCUU	1740	AAGCATTC T GAGCTGGC	2464
2978	AACAAGCC CUGAUGAG X CGAA ICUCAGAA	1741	TTCTGAGC T GGCTTGTT	2465
2982	AAACAACA CUGAUGAG X CGAA ICCAGCUC	1742	GAGCTGGC T TGTGTTT	2466
2998	CAUAUACA CUGAUGAG X CGAA IACUUAUA	1743	TTTAAGTC C TGTATATG	2467
2999	ACAUUAC CUGAUGAG X CGAA IGACUUA	1744	TTAAGTCC T GTATATGT	2468
3040	UUUUGAAA CUGAUGAG X CGAA ICUACUAU	1745	ATAGTAGC A TTTCAAAA	2469
3045	GUCCAUUU CUGAUGAG X CGAA IAAAUAGCU	1746	AGCATTTTC A AAATGGAC	2470
3058	GUUAAACC CUGAUGAG X CGAA IUACGUCC	1747	GGACGTAC T GGTTTAAC	2471
3067	GGAUAGGA CUGAUGAG X CGAA IUUAAACC	1748	GGTTTAAC C TCCTATCC	2472
3068	AGGAUAGG CUGAUGAG X CGAA IGUUAAC	1749	GTTTAACC T CCTATCCT	2473
3070	CAAGGAUA CUGAUGAG X CGAA IAGGUUA	1750	TTAACCTC C TATCCTTG	2474
3071	CCAAGGAU CUGAUGAG X CGAA IGAGGUUA	1751	TAACCTCC T ATCCTTGG	2475
3075	CUCUCCAA CUGAUGAG X CGAA IAUAGGAG	1752	CTCCTATC C TTGGAGAG	2476
3076	GCUCUCCA CUGAUGAG X CGAA IGAUAGGA	1753	TCCTATCC T TGGAGAGC	2477
3085	GAGCCAGC CUGAUGAG X CGAA ICUCUCCA	1754	TGGAGAGC A GCTGGCTC	2478
3088	GGAGAGCC CUGAUGAG X CGAA ICUGCUCU	1755	AGAGCAGC T GGCTCTCC	2479
3092	AGGUGGAG CUGAUGAG X CGAA ICCAGCUG	1756	CAGCTGGC T CTCCACCT	2480
3094	CAAGGUGG CUGAUGAG X CGAA IAGCCAGC	1757	GCTGGCTC T CCACCTTG	2481

Table 4

3096	AACAAGGU CUGAUGAG X CGAA IAGAGCCA	1758	TGGCTCTC C ACCTTGTT	2482
3097	UAACAAGG CUGAUGAG X CGAA IGAGAGCC	1759	GGCTCTCC A CCTTGTTA	2483
3099	UGUAACAA CUGAUGAG X CGAA IUGGAGAG	1760	CTCTCCAC C TTGTTACA	2484
3100	GUGUAACA CUGAUGAG X CGAA IGUGGAGA	1761	TCTCCACC T TGTACAC	2485
3107	ACAUAUUG CUGAUGAG X CGAA IUAACAAG	1762	CTTGTTAC A CATTATGT	2486
3109	UAACAUAA CUGAUGAG X CGAA IUGUAACA	1763	TGTTACAC A TTATGTTA	2487
3132	AGCAGAGC CUGAUGAG X CGAA ICUCGCUA	1764	TAGCGAGC T GCTCTGCT	2488
3135	CAUAGCAG CUGAUGAG X CGAA ICAGCUCG	1765	CGAGCTGC T CTGCTATG	2489
3137	GACAUAGC CUGAUGAG X CGAA IAGCAGCU	1766	AGCTGCTC T GCTATGTC	2490
3140	AAGGACAU CUGAUGAG X CGAA ICAGAGCA	1767	TGCTCTGC T ATGTCCTT	2491
3146	UGGCUUAA CUGAUGAG X CGAA IACAUAGC	1768	GCTATGTC C TTAAGCCA	2492
3147	UUGGCUUA CUGAUGAG X CGAA IGACAUAG	1769	CTATGTCC T TAAGCCAA	2493
3153	UAAAUUUU CUGAUGAG X CGAA ICUUUAGG	1770	CCTTAAGC C AATATTTA	2494
3154	GUAAUAU CUGAUGAG X CGAA IGCUUUAG	1771	CTTAAGCC A ATATTTAC	2495
3163	ACCUGAUG CUGAUGAG X CGAA IUAAUAU	1772	ATATTTAC T CATCAGGT	2496
3165	UGACCUGA CUGAUGAG X CGAA IAGUAAU	1773	ATTTACTC A TCAGGTCA	2497
3168	UAAUGACC CUGAUGAG X CGAA IAUGAGUA	1774	TACTCATC A GGTCATTA	2498
3173	AAAAUAA CUGAUGAG X CGAA IACCUGAU	1775	ATCAGGTC A TTATTTT	2499
3185	AUGGCCAU CUGAUGAG X CGAA IUAAAAA	1776	TTTTTTAC A ATGGCCAT	2500
3191	UAUCCAU CUGAUGAG X CGAA ICCAUUGU	1777	ACAATGGC C ATGGAATA	2501
3192	UUAUCCA CUGAUGAG X CGAA IGCCAUUG	1778	CAATGGCC A TGGAATAA	2502
3203	GUAAAAU CUGAUGAG X CGAA IUUUUUC	1779	GAATAAAC C ATTTTAC	2503
3204	UGUAAAA CUGAUGAG X CGAA IGUUUAU	1780	AATAAACC A TTTTACA	2504

Table 5

Table 5: Human PTP-1B G-Cleaver Ribozyme and Target Sequence

Nt. position	Ribozyme Sequence	Seq. ID Nos.	Substrate Sequence	Seq. ID Nos.
25	CACUG UGAUG GCAUGCACUAUGC GCG GCCGUCUGCC	2505	GGCAGACGGC G CAGTG	2670
35	CUUCU UGAUG GCAUGCACUAUGC GCG GGCCACACUGC	2506	GCAGTGGGCC G AGAAG	2671
46	UGCUG UGAUG GCAUGCACUAUGC GCG GCCUCCUUCU	2507	AGAAGGAGGC G CAGCA	2672
55	CAGGG UGAUG GCAUGCACUAUGC GCG GCGUCGUGCG	2508	CGCAGCAGCC G CCCTG	2673
89	CUGCU UGAUG GCAUGCACUAUGC GCG GAACUCCUUCU	2509	AAAGGAGTTC G AGCAG	2674
98	CUUGU UGAUG GCAUGCACUAUGC GCG GAUCUGCUCG	2510	CGAGCAGATC G ACAAG	2675
138	CAUGU UGAUG GCAUGCACUAUGC GCG GGAUAUCCUG	2511	CAGGATATCC G ACATG	2676
143	GGCUU UGAUG GCAUGCACUAUGC GCG AUGUGGGAUA	2512	TATCCGACAT G AAGCC	2677
152	GAAGU UGAUG GCAUGCACUAUGC GCG ACUGGCUUCA	2513	TGAAGCCAGT G ACTTC	2678
195	UAUUU UGAUG GCAUGCACUAUGC GCG GGUUUUUGUU	2514	AACAAAACC G AAATA	2679
224	AUGGU UGAUG GCAUGCACUAUGC GCG AAAGGGACUG	2515	CAGTCCTTT G ACCAT	2680
260	AUAGU UGAUG GCAUGCACUAUGC GCG AUUAUCUUUCU	2516	AGAAGATAAT G ACTAT	2681
272	ACUAG UGAUG GCAUGCACUAUGC GCG GUUGAUUAAG	2517	CTATATCAAC G CTAGT	2682
280	UUUAU UGAUG GCAUGCACUAUGC GCG AAACUAGCGU	2518	ACGCTAGTTT G ATAAA	2683
331	UUAGG UGAUG GCAUGCACUAUGC GCG AAAGGGCCCU	2519	AGGGCCCTTT G CCTAA	2684
342	GACCG UGAUG GCAUGCACUAUGC GCG AUGUGUUAGG	2520	CCTAACACAT G CGGTC	2685
394	UUAG UGAUG GCAUGCACUAUGC GCG AUGACGACAC	2521	GTGTCGTCAT G CTCAA	2686
406	UCCAU UGAUG GCAUGCACUAUGC GCG ACUCUGUUUA	2522	TCAACAGAGT G ATGGA	2687
429	GUGCG UGAUG GCAUGCACUAUGC GCG AUUUUAAAGA	2523	TCGTTAAAAT G CGCAC	2688
431	UUGUG UGAUG GCAUGCACUAUGC GCG GCAUUUUAAAC	2524	GTTAAAATGC G CACAA	2689
466	AAGAU UGAUG GCAUGCACUAUGC GCG AUCUCUUUUU	2525	AAAAAGAGAT G ATCTT	2690
473	GUCUU UGAUG GCAUGCACUAUGC GCG AAAGAUAUUC	2526	GATGATCTTT G AAGAC	2691
487	AAUUU UGAUG GCAUGCACUAUGC GCG AAUUUUUGUU	2527	ACACAAATTT G AAATT	2692
499	GAGAU UGAUG GCAUGCACUAUGC GCG AAUGUUAAUU	2528	AATTAAACAT G ATCTC	2693
506	AUCUU UGAUG GCAUGCACUAUGC GCG AGAGAUCAAU	2529	ATTGATCTCT G AAGAT	2694
532	UGUCG UGAUG GCAUGCACUAUGC GCG ACUGUAUAUU	2530	ATTATACAGT G CGACA	2695
534	GCUGU UGAUG GCAUGCACUAUGC GCG GCACUGUAUA	2531	TATACAGTGC G ACAGC	2696
573	UCUCU UGAUG GCAUGCACUAUGC GCG GAGUUUCUUG	2532	CAAGAAACTC G AGAGA	2697

Table 5

608	AAAGU UGAUG GCAUGCACUAUGC GCG AGGCCAUGUG	2533	CACATGGCCT G ACTTT	2698
623	UGAUU UGAUG GCAUGCACUAUGC GCG AGGGACUCCA	2534	TGGAGTCCCT G AATCA	2699
643	AAGUU UGAUG GCAUGCACUAUGC GCG AAGAAUGAGG	2535	CCTCATTTCTT G AACTT	2700
663	ACUCU UGAUG GCAUGCACUAUGC GCG GGACUUUGAA	2536	TTCAAAGTCC G AGAGT	2701
706	CAGUG UGAUG GCAUGCACUAUGC GCG ACCACAACGG	2537	CCGTGTGGT G CACTG	2702
711	CACUG UGAUG GCAUGCACUAUGC GCG AGUGCACAC	2538	GTGGTGCACT G CAGTG	2703
716	GCCUG UGAUG GCAUGCACUAUGC GCG ACUGCAGUGC	2539	GCACTGCAGT G CAGGC	2704
752	GGUAU UGAUG GCAUGCACUAUGC GCG AGCCAGACAG	2540	CTGTCTGGCT G ATACC	2705
759	AGAGG UGAUG GCAUGCACUAUGC GCG AGGUUACAGC	2541	GCTGATACCT G CCTCT	2706
766	AUCAG UGAUG GCAUGCACUAUGC GCG AAGAGGCAGG	2542	CCTGCCTCTT G CTGAT	2707
769	UCCAU UGAUG GCAUGCACUAUGC GCG AGCAAGAGGC	2543	GCCTCTTGCT G ATGGA	2708
800	GAUUA UGAUG GCAUGCACUAUGC GCG AACGGAAGAA	2544	TTCTTCCGTT G ATATC	2709
814	AACAG UGAUG GCAUGCACUAUGC GCG ACUUUCUUGA	2545	TCAAGAAAGT G CTGTT	2710
826	UCCU UGAUG GCAUGCACUAUGC GCG AUUUCUAACA	2546	TGTTAGAAAT G AGGAA	2711
847	UGGAU UGAUG GCAUGCACUAUGC GCG AGCCCCAUCC	2547	GGATGGGGCT G ATCCA	2712
860	CUGGU UGAUG GCAUGCACUAUGC GCG GGCUGUCUGG	2548	CCAGACAGCC G ACCAG	2713
868	AAGCG UGAUG GCAUGCACUAUGC GCG AGCUGGUCGG	2549	CCGACCAGCT G CGCTT	2714
870	AGAAG UGAUG GCAUGCACUAUGC GCG GCAGCUGGUC	2550	GACCAGCTGC G CTCT	2715
889	UCGAU UGAUG GCAUGCACUAUGC GCG ACAGCCAGGU	2551	ACCTGGCTGT G ATCGA	2716
893	ACCUU UGAUG GCAUGCACUAUGC GCG GAUCACAGCC	2552	GGCTGTGATC G AAGGT	2717
899	UUUGG UGAUG GCAUGCACUAUGC GCG ACCUUCGAUC	2553	GATCGAAGGT G CCAAA	2718
928	UCCUG UGAUG GCAUGCACUAUGC GCG ACGGAAGAGU	2554	ACTCTTCCGT G CAGGA	2719
956	GUCCU UGAUG GCAUGCACUAUGC GCG GUGGGAAGC	2555	GCTTTCCAC G AGGAC	2720
977	AUGCU UGAUG GCAUGCACUAUGC GCG GGGUGGGGC	2556	GCCCCACCC G AGCAT	2721
1011	GGAUU UGAUG GCAUGCACUAUGC GCG GUUUGGGUGG	2557	CCACCCAAAC G AATCC	2722
1038	CCCUG UGAUG GCAUGCACUAUGC GCG AUUUCUUAU	2558	AATGGGAAAT G CAGGG	2723
1069	UCCUU UGAUG GCAUGCACUAUGC GCG ACCACUGGU	2559	ACCAGTGGT G AAGGA	2724
1098	UGGGG UGAUG GCAUGCACUAUGC GCG AGUUCUUUUC	2560	GATAAAGACT G CCCCC	2725
1133	UGCGG UGAUG GCAUGCACUAUGC GCG AUUUAAGGGG	2561	CCCCTTAAAT G CCGCA	2726
1136	GGGUG UGAUG GCAUGCACUAUGC GCG GGCAUUUAAG	2562	CTTAAATGCC G CACCC	2727
1151	GCUUU UGAUG GCAUGCACUAUGC GCG GAUGCCGUAG	2563	CTACGGCATC G AAAGC	2728

Table 5

1159	UGACU UGAUG GCAUGCACUAUGC GCG AUGCUUUCGA	2564	TCGAAAGCAT G AGTCA	2729
1172	AACUU UGAUG GCAUGCACUAUGC GCG AGUGUCUUGA	2565	TCAAGACACT G AAGTT	2730
1206	CACCU UGAUG GCAUGCACUAUGC GCG GAAGACUUC	2566	GGAAGTCTTC G AGGTG	2731
1211	CUGGG UGAUG GCAUGCACUAUGC GCG ACCUCGAAGA	2567	TCTTCGAGGT G CCAG	2732
1220	GGAGG UGAUG GCAUGCACUAUGC GCG AGCCUGGGCA	2568	TGCCCAGGCT G CCTCC	2733
1249	UGGGG UGAUG GCAUGCACUAUGC GCG AGUGACGGCU	2569	AGCGGTCACT G CCGA	2734
1253	CUUCU UGAUG GCAUGCACUAUGC GCG GGGCAGUGAC	2570	GTCACTGCCC G AGAAG	2735
1262	GUCCU UGAUG GCAUGCACUAUGC GCG GUCCUUCUG	2571	CGAGAAGGAC G AGGAC	2736
1271	CAGUG UGAUG GCAUGCACUAUGC GCG AUGGUCCUCG	2572	CGAGGACCAT G CACTG	2737
1276	UAACU UGAUG GCAUGCACUAUGC GCG AGUGCAUGGU	2573	ACCATGCACT G AGTTA	2738
1308	CCACG UGAUG GCAUGCACUAUGC GCG ACAUGUUGAC	2574	GTCAACATGT G CGTGG	2739
1334	GUAAAG UGAUG GCAUGCACUAUGC GCG GCCGGCCGUG	2575	CACGGCCGGC G CTTAC	2740
1344	UGUAG UGAUG GCAUGCACUAUGC GCG AGAGGUAAGC	2576	GCTTACCTCT G CTACA	2741
1379	AGGGU UGAUG GCAUGCACUAUGC GCG AGGCUAUGUG	2577	CACATAGCCT G ACCCT	2742
1412	AGAGG UGAUG GCAUGCACUAUGC GCG GGACAGUGGG	2578	CCCAGTGTCC G CCTCT	2743
1418	GCGGG UGAUG GCAUGCACUAUGC GCG AGAGGCGGAC	2579	GTCCGCTCT G CCCGC	2744
1422	CUUCU UGAUG GCAUGCACUAUGC GCG GGGCAGAGGC	2580	GCCTCTGCC G CAGAG	2745
1433	UCGGG UGAUG GCAUGCACUAUGC GCG GUGGUCUCUG	2581	CAGAGCCAC G CCCGA	2746
1437	CUAGU UGAUG GCAUGCACUAUGC GCG GGGCGUGGGC	2582	GCCACGCCC G ACTAG	2747
1450	CGCGG UGAUG GCAUGCACUAUGC GCG AUGCCUGCUA	2583	TAGCAGGCAT G CCGCG	2748
1453	UACCG UGAUG GCAUGCACUAUGC GCG GGCAUGCCUG	2584	CAGGCATGCC G CGGTA	2749
1469	UCCGG UGAUG GCAUGCACUAUGC GCG GGCCCUUACC	2585	GGTAAGGGCC G CCGGA	2750
1477	CUACG UGAUG GCAUGCACUAUGC GCG GGUCCGGCGG	2586	CGCCGGACC G CGTAG	2751
1513	UAGUG UGAUG GCAUGCACUAUGC GCG AGAACCAACG	2587	CGTTGGTTCT G CACTA	2752
1569	AGGGG UGAUG GCAUGCACUAUGC GCG AAAAGUAAA	2588	TTTACTTTT G CCCCT	2753
1583	GUACU UGAUG GCAUGCACUAUGC GCG AAAGUGGAAG	2589	CTTCCACTTT G AGTAC	2754
1610	CUCCU UGAUG GCAUGCACUAUGC GCG AAAAAUUGGC	2590	GCCATTTTT G AGGAG	2755
1619	UCUUU UGAUG GCAUGCACUAUGC GCG ACUCUCCUCA	2591	TGAGGAGAGT G AAAGA	2756
1634	GCCAG UGAUG GCAUGCACUAUGC GCG AUGGUACUCU	2592	AGAGTACCAT G CTGGC	2757
1643	CUCUG UGAUG GCAUGCACUAUGC GCG GCCGCCAGCA	2593	TGCTGGCGGC G CAGAG	2758
1678	UGGGG UGAUG GCAUGCACUAUGC GCG GAGCCCCAAG	2594	CTTGGGGCTC G CCCCCA	2759

Table 5

1723	UCUG UGAUG GCAUGCACUAUGC GCG GUGCGCGCCG	2595	CGGCGGGCAC G CCAGA	2760
1742	AGAUU UGAUG GCAUGCACUAUGC GCG AAGGGGGG	2596	CCCCCCCCCTT G AATCT	2761
1748	CCUG UGAUG GCAUGCACUAUGC GCG AGAUCAAGG	2597	CCTTGAATCT G CAGGG	2762
1811	UAGUG UGAUG GCAUGCACUAUGC GCG ACUAUGGAG	2598	CATCCATAGT G CACTA	2763
1827	UGGUU UGAUG GCAUGCACUAUGC GCG AAGAAAUGC	2599	GCAATTTCTT G AACCA	2764
1853	GACAU UGAUG GCAUGCACUAUGC GCG AAAAAUUU	2600	AAAATTTTT G ATGTC	2765
1865	UGAUG UGAUG GCAUGCACUAUGC GCG AAGGUGACA	2601	TGTCAGCCTT G CATCA	2766
1931	CAUGG UGAUG GCAUGCACUAUGC GCG AAGGCUUCC	2602	GGAAGGCTTT G CCATG	2767
1942	CGCAG UGAUG GCAUGCACUAUGC GCG AGGCCAUGG	2603	CCATGGGCTT G CTGCG	2768
1945	UGACG UGAUG GCAUGCACUAUGC GCG AGCAGGCCCA	2604	TGGGCGCTGCT G CGTCA	2769
1997	AAUAU UGAUG GCAUGCACUAUGC GCG ACUAAAUAA	2605	GTATTTAGT G ATATT	2770
2014	CUUCU UGAUG GCAUGCACUAUGC GCG ACGUACCCA	2606	TGGGTAACGT G AGAAG	2771
2030	UAUAG UGAUG GCAUGCACUAUGC GCG AUGGUUCUUA	2607	ATAGAACAAT G CTATA	2772
2045	GUGUU UGAUG GCAUGCACUAUGC GCG AUUAUAUUA	2608	AATATATAAT G AACAC	2773
2073	CACAU UGAUG GCAUGCACUAUGC GCG AUGUUUCUUA	2609	TAAGAAACAT G ATGTG	2774
2078	AAUCU UGAUG GCAUGCACUAUGC GCG ACAUCAUGUU	2610	AACATGATGT G AGATT	2775
2094	AUAAG UGAUG GCAUGCACUAUGC GCG GGGACAAAGU	2611	ACTTTGTCCC G CTTAT	2776
2103	GGGAG UGAUG GCAUGCACUAUGC GCG AGAAUAAAGC	2612	CGCTTATTTCT G CTCCC	2777
2117	UCUAG UGAUG GCAUGCACUAUGC GCG AGAAUAAAGC	2613	CCTGTTATCT G CTAGA	2778
2140	GGGAG UGAUG GCAUGCACUAUGC GCG AGUUAUAG	2614	CTCAATCACT G CTCCC	2779
2162	ACAUG UGAUG GCAUGCACUAUGC GCG AUUCUAAUAC	2615	GTATTAGAT G CATGT	2780
2186	UUCAU UGAUG GCAUGCACUAUGC GCG AGGACACAAG	2616	CTTGTCCTT G ATGAA	2781
2189	UUUUU UGAUG GCAUGCACUAUGC GCG AUCAGGACAC	2617	GTGTCCTGAT G AAAA	2782
2200	UCAAG UGAUG GCAUGCACUAUGC GCG ACAUAAUUUU	2618	AAAAATATGT G CTTGA	2783
2204	CAUUU UGAUG GCAUGCACUAUGC GCG AAGCACAUAU	2619	ATATGTGCTT G AATG	2784
2209	UUUCU UGAUG GCAUGCACUAUGC GCG AUUUAAGCA	2620	TGCTTGAAAT G AGAAA	2785
2219	GAGAU UGAUG GCAUGCACUAUGC GCG AAAGUUUUC	2621	GAGAACTTT G ATCTC	2786
2226	GUAAU UGAUG GCAUGCACUAUGC GCG AGAGAUCAAA	2622	TTTGATCTCT G CTTAC	2787
2238	UGGGG UGAUG GCAUGCACUAUGC GCG ACAUUAAGUA	2623	TTACTAATGT G CCCCC	2788
2260	ACAGG UGAUG GCAUGCACUAUGC GCG AGGUUGGACU	2624	AGTCCACCT G CCTGT	2789
2266	UCAUG UGAUG GCAUGCACUAUGC GCG ACAGGCAGGU	2625	ACCTGCCTGT G CATGA	2790

Table 5

2270	CAGGU UGAUG GCAUGCACUAUGC GCG AUGCACAGGC	2626	GCCTGTGCAT G ACCTG	2791
2275	AUGAU UGAUG GCAUGCACUAUGC GCG AGGUCAUGCA	2627	TGCATGACCT G ATCAT	2792
2308	UUCAG UGAUG GCAUGCACUAUGC GCG AACAGGCUUA	2628	TAAGCCTGTT G CTGAA	2793
2311	GACUU UGAUG GCAUGCACUAUGC GCG AGCAACAGGC	2629	GCCTGTTGCT G AAGTC	2794
2323	CUGAG UGAUG GCAUGCACUAUGC GCG GACAAUGACU	2630	AGTCATTGTC G CTCAG	2795
2338	AACUG UGAUG GCAUGCACUAUGC GCG ACCCUAUGC	2631	GCAATAGGGT G CAGTT	2796
2362	UUAGG UGAUG GCAUGCACUAUGC GCG AAUUGCCUUA	2632	ATAGGCATT G CCTAA	2797
2378	AGUGU UGAUG GCAUGCACUAUGC GCG AUGCCAGGAA	2633	TTCCTGGCAT G ACACT	2798
2389	GAAGU UGAUG GCAUGCACUAUGC GCG ACUAGAGUGU	2634	ACACTCTAGT G ACTTC	2799
2400	GGCCU UGAUG GCAUGCACUAUGC GCG ACCAGGAAGU	2635	ACTTCTGGT G AGGCC	2800
2433	UACAG UGAUG GCAUGCACUAUGC GCG AAGACCCUGC	2636	GCAGGGTCTT G CTGTA	2801
2483	CAGAG UGAUG GCAUGCACUAUGC GCG GUGAGGUGUG	2637	CACACCTCAC G CTCTG	2802
2494	UAAAU UGAUG GCAUGCACUAUGC GCG AUGUCCAGAG	2638	CTCTGGACAT G ATTTA	2803
2520	GGGGG UGAUG GCAUGCACUAUGC GCG GGGGGGUGUC	2639	GACACCCCCC G CCCCC	2804
2546	AAUGG UGAUG GCAUGCACUAUGC GCG GGAGGCUGAU	2640	ATCAGCCTCC G CCATT	2805
2559	AGUGU UGAUG GCAUGCACUAUGC GCG GACUUGGAUU	2641	ATTCCAAGTC G ACACT	2806
2571	CUGCU UGAUG GCAUGCACUAUGC GCG AGAAGAGUG	2642	CACCTTCTT G AGCAG	2807
2582	CAAAU UGAUG GCAUGCACUAUGC GCG ACGUUCUGCU	2643	AGCAGACCGT G ATTTG	2808
2602	UCCAG UGAUG GCAUGCACUAUGC GCG AGGUGCCUCU	2644	AGAGGCACCT G CTGGA	2809
2621	UGUUU UGAUG GCAUGCACUAUGC GCG AAGAAUGUG	2645	CACACTTCTT G AAACA	2810
2635	ACCGU UGAUG GCAUGCACUAUGC GCG ACCAGGUCUG	2646	CAGCCTGGGT G ACGGT	2811
2655	GGCGG UGAUG GCAUGCACUAUGC GCG AGGUGCCCUA	2647	TAGGCAGCCT G CCGCC	2812
2658	GACGG UGAUG GCAUGCACUAUGC GCG GGCAAGGUGA	2648	GCAGCCTGCC G CCGTC	2813
2682	CUGGG UGAUG GCAUGCACUAUGC GCG AAGGUGAACC	2649	GGTTCACTT G CCGAG	2814
2685	UCUCU UGAUG GCAUGCACUAUGC GCG GCGUUCUCUG	2650	TCACCTTGCC G AGAGA	2815
2694	AGACG UGAUG GCAUGCACUAUGC GCG GCCUCUCUCG	2651	CGAGAGAGGC G CGTCT	2816
2700	UGGGG UGAUG GCAUGCACUAUGC GCG AGACGGGCUU	2652	AGGCGGTCT G CCCC	2817
2727	ACCAU UGAUG GCAUGCACUAUGC GCG AGGCCCCACA	2653	TGTGGGCCT G ATGGT	2818
2733	GUGAG UGAUG GCAUGCACUAUGC GCG ACCAUCAGGC	2654	GCCTGATGGT G CTCAC	2819
2739	AGAGU UGAUG GCAUGCACUAUGC GCG GUGAGCACCA	2655	TGGTGCTCAC G ACTCT	2820
2749	CUUUG UGAUG GCAUGCACUAUGC GCG AGGAAGAGUC	2656	GACTCTTCTT G CAAAG	2821

Table 5

2761	GUCUU	UGAUG	GCAUGCACUAUGC	GCG	AGUUCUCCUU	2657	AAAGGGAAC	G	AAGAC	2822
2793	UUUUU	UGAUG	GCAUGCACUAUGC	GCG	AUGUUA AAAA	2658	TTTTTAACAT	G	AAAAA	2823
2818	UAGCU	UGAUG	GCAUGCACUAUGC	GCG	GGGAGCUACA	2659	TGTAGCTCCC	G	AGCTA	2824
2831	GCUGG	UGAUG	GCAUGCACUAUGC	GCG	AAGAGAGUAG	2660	CTACTCTCTT	G	CCAGC	2825
2850	AAAGG	UGAUG	GCAUGCACUAUGC	GCG	AAAUGUGAA	2661	TTCACATTTT	G	CCTTT	2826
2902	CACCU	UGAUG	GCAUGCACUAUGC	GCG	GAUUGUCC	2662	GGGAACATTC	G	AGGTG	2827
2915	CUCUG	UGAUG	GCAUGCACUAUGC	GCG	AGGUGACAC	2663	GTGTCACCC	G	CAGAG	2828
2928	CACCU	UGAUG	GCAUGCACUAUGC	GCG	ACCAUAGCUC	2664	GAGCTATGGT	G	AGGTG	2829
2950	CCUGG	UGAUG	GCAUGCACUAUGC	GCG	ACCUAAGCCU	2665	AGGCTTAGGT	G	CCAGG	2830
2969	CAGCU	UGAUG	GCAUGCACUAUGC	GCG	AGAAUGCUUA	2666	TAAGCATTTCT	G	AGCTG	2831
3123	CAGCU	UGAUG	GCAUGCACUAUGC	GCG	GCUACCCUC	2667	GAGAGGTAGC	G	AGCTG	2832
3128	CAGAG	UGAUG	GCAUGCACUAUGC	GCG	AGCUCCCUAC	2668	GTAGCGAGCT	G	CTCTG	2833
3133	CAUAG	UGAUG	GCAUGCACUAUGC	GCG	AGAGCAGCUC	2669	GAGCTGCTCT	G	CTATG	2834

Table 6

Table 6: Human PTP-1B DNAzyme and Target Sequence

Nt. Position	DNAzyme Sequence	Seq. ID Nos.	Substrate Sequence	Seq. ID Nos.
11	GCTCTAGG GGCTAGCTACAACGA CGCGTGCG	2835	GCACGCG G CCUAGAGC	3545
18	GTCTGCCG GGCTAGCTACAACGA TCTAGGCC	2836	GGCCUAGA G CGGCAGAC	3546
21	GCGTCTG GGCTAGCTACAACGA CGCTCTAG	2837	CUAGAGCG G CAGACGGC	3547
25	CTGCGCCG GGCTAGCTACAACGA CTGCGGCT	2838	AGCGGCAG A CGGCGCAG	3548
28	CCACTGCG GGCTAGCTACAACGA CGTCTGCC	2839	GGCAGACG G CGCAGUGG	3549
30	GCCCACTG GGCTAGCTACAACGA GCCGTCTG	2840	CAGACGGC G CAGUGGGC	3550
33	TGGGCCCA GGCTAGCTACAACGA TGGCCGCT	2841	ACGGCGCA G UGGGCCGA	3551
37	CTTCTCGG GGCTAGCTACAACGA CCACTGCG	2842	CGCAGUGG G CCGAGAAG	3552
49	CTGCTGCG GGCTAGCTACAACGA CTCCTTCT	2843	AGAAGGAG G CGCAGCAG	3553
51	GGCTGCTG GGCTAGCTACAACGA GCCTCCTT	2844	AAGGAGGC G CAGCAGCC	3554
54	GGGGGCTG GGCTAGCTACAACGA TGGCCCTC	2845	GAGGCGCA G CAGCCGCC	3555
57	CAGGGCGG GGCTAGCTACAACGA TGCTGCGC	2846	GCGCAGCA G CCGCCCTG	3556
60	GGCCAGGG GGCTAGCTACAACGA GGCTGCTG	2847	CAGCAGCC G CCCUGGCC	3557
66	ATGACGGG GGCTAGCTACAACGA CAGGGCGG	2848	CCGCCCTG G CCCGUCAU	3558
70	CTCCATGA GGCTAGCTACAACGA GGGCCAGG	2849	CTUGGCC G UCAUGGAG	3559
73	CATCTCCA GGCTAGCTACAACGA GACGGGCC	2850	GGCCCGUC A UGGAGAUG	3560
79	CTTTTCCA GGCTAGCTACAACGA CTCCATGA	2851	UCAUGGAG A UGGAAAAG	3561
90	TGCTCGAA GGCTAGCTACAACGA TCCTTTTC	2852	GAAAAGGA G UUCGAGCA	3562
96	TCGATCTG GGCTAGCTACAACGA TCGAATCT	2853	GAGUUCGA G CAGAUCGA	3563
100	CTTGTCGA GGCTAGCTACAACGA CTGCTCGA	2854	UCGAGCAG A UCGACAAG	3564
104	CGGACTTG GGCTAGCTACAACGA CGATCTGC	2855	GCAGAUCC A CAAGUCCG	3565
108	CTCCCGGA GGCTAGCTACAACGA TTGTCGAT	2856	AUCGACAA G UCCGGGAG	3566
116	CGGCCCGG GGCTAGCTACAACGA TCCCGGAC	2857	GUCCGGGA G CUGGGCGG	3567
121	AATGGCCG GGCTAGCTACAACGA CCAGCTCC	2858	GGAGCUGG G CGGCCAUU	3568
124	GTAATGGG GGCTAGCTACAACGA GCGCCAGC	2859	GCUGGGCG G CCAUUUAC	3569
127	CTGGTAAA GGCTAGCTACAACGA GGCCGCCC	2860	GGCGGCC A UUUACCAG	3570
131	TATCCTGG GGCTAGCTACAACGA AAATGGCC	2861	GGCCAUUU A CCAGGAUA	3571
137	GTGGGATA GGCTAGCTACAACGA CCTGGTAA	2862	UUACCAGG A UAUCCGAC	3572

Table 6

139	ATGTCGGA GGCTAGCTACAACGA ATCCTGGT	2863	ACCAGGAU A UCCGACAU	3573
144	GCTTCATG GGCTAGCTACAACGA CGGATATC	2864	GAUAUCCG A CAUGAAGC	3574
146	TGGCTTCA GGCTAGCTACAACGA GTCGGATA	2865	UAUCCGAC A UGAAGCCA	3575
151	GTCACTGG GGCTAGCTACAACGA TTCATGTC	2866	GACAUAG A G CAGUGAC	3576
155	GGAAGTCA GGCTAGCTACAACGA TGGCTTCA	2867	UGAAGCCA G UGACUUC	3577
158	ATGGGAAG GGCTAGCTACAACGA CACTGGCT	2868	AGCAGUG A CUUCCCAU	3578
165	ACTCTACA GGCTAGCTACAACGA GGGAAATC	2869	GACUCCCC A UGUAGAGU	3579
167	CCACTCTA GGCTAGCTACAACGA ATGGGAAG	2870	CUUCCCAU G UAGUGUG	3580
172	CTTGGCCA GGCTAGCTACAACGA TCTACATG	2871	CAUGUAG A UGGCCNAG	3581
175	AAGCTTGG GGCTAGCTACAACGA CACTCTAC	2872	GUAGAGUG G CCAAGCUU	3582
180	TTAGGAAG GGCTAGCTACAACGA TTGGCCAC	2873	GUGGCCAA G CUUCCUAA	3583
191	GGTTTTTG GGCTAGCTACAACGA TCTTAGGA	2874	UCCUAGA A CAAAACC	3584
197	TATTTCCG GGCTAGCTACAACGA TTTTGTTC	2875	GAACAAA A CCGAAUA	3585
203	TGTACCTA GGCTAGCTACAACGA TTCGGTTT	2876	AAACCGAA A UAGGUACA	3586
207	TCTCTGTA GGCTAGCTACAACGA CTATTTCC	2877	CGAAUAG G UACAGAGA	3587
209	CGTCTCTG GGCTAGCTACAACGA ACTATTTT	2878	AAUAGGU A CAGAGAGC	3588
215	GACTGACG GGCTAGCTACAACGA CTCTGTAC	2879	GUACAGAG A CGUCAGUC	3589
217	GGGACTGA GGCTAGCTACAACGA GTCTCTGT	2880	ACAGAGAC G UCAGUCCC	3590
221	CAAAGGGA GGCTAGCTACAACGA TGACGTCT	2881	AGACGUCA G UCCCUIUG	3591
230	GACTATGG GGCTAGCTACAACGA CAAAGGGA	2882	UCCCUUUG A CCAUAGUC	3592
233	TCCGACTA GGCTAGCTACAACGA GGTCAAAG	2883	CUUUGACC A UAGUCGGA	3593
236	TAATCCGA GGCTAGCTACAACGA TATGGTCA	2884	UGACCAUA G UCGGAUUA	3594
241	TAGTTTAA GGCTAGCTACAACGA CCGACTAT	2885	AUAGUGGG A UUAACCUA	3595
246	TGATGTAG GGCTAGCTACAACGA TTAATCCG	2886	CGGAUJAA A CUACAUA	3596
249	TCTTGATG GGCTAGCTACAACGA AGTTTAAT	2887	AUUAACU A CAUCAAGA	3597
251	CTTCTTGA GGCTAGCTACAACGA GTAGTTTA	2888	UAAACUAC A UCAAGNAG	3598
260	AGTCATTA GGCTAGCTACAACGA CTTCTTGA	2889	UCAAGNAG A UAAUGACU	3599
263	TATAGTCA GGCTAGCTACAACGA TATCTTCT	2890	AGAAGUA A UGACUUA	3600
266	TGATATAG GGCTAGCTACAACGA CATTATCT	2891	AGAUAUG A CUUAUUA	3601
269	CGTTGATA GGCTAGCTACAACGA AGTCATTA	2892	UAAUGACU A UAUCAACG	3602
271	AGCGTTGA GGCTAGCTACAACGA ATAGTCAT	2893	AUGACUUA A UCAACGCU	3603

Table 6

275	AACTAGCG	GGCTAGCTACAACGA	TGATATAG	2894	CUAUAUCA	A CGUAGUU	3604
277	CAAACTAG	GGCTAGCTACAACGA	GTTGATAT	2895	AUAUCAAC	G CUAGUUUG	3605
281	TTATCAAA	GGCTAGCTACAACGA	TAGCGTTG	2896	CAACGCUA	G UUGAUAA	3606
286	CATTTTAA	GGCTAGCTACAACGA	CAAACTAG	2897	CUAGUUUG	A UAAAAUG	3607
292	TTCTTCCA	GGCTAGCTACAACGA	TTTTATCA	2898	UGAUAAAA	A UGGAAGAA	3608
301	CCTTTGGG	GGCTAGCTACAACGA	TTCTTCCA	2899	UGGAAGAA	G CCCAAGG	3609
311	GAATGTAA	GGCTAGCTACAACGA	TCCTTTGG	2900	CCAAAGGA	G UUAUAUUC	3610
314	TAAGAATG	GGCTAGCTACAACGA	AACTCCTT	2901	AAGGAGUU	A CAUUCUUA	3611
316	GGTAAGAA	GGCTAGCTACAACGA	GTAATCTC	2902	GGAGUUAC	A UUCUUACC	3612
322	GCCCTGGG	GGCTAGCTACAACGA	AAGAAATG	2903	ACAUUCUU	A CCCAGGGC	3613
329	GCAAGGGG	GGCTAGCTACAACGA	CCTGGGTA	2904	UACCCAGG	G CCCUUUGC	3614
336	GTGTTAGG	GGCTAGCTACAACGA	AAAGGGCC	2905	GGCCCCUU	G CGUAACAC	3615
341	CGCATGTG	GGCTAGCTACAACGA	TAGGCAAA	2906	UUUGCCUA	A CACAUGCG	3616
343	ACCGCATG	GGCTAGCTACAACGA	GTTAGGCA	2907	UGCCUAAC	A CAUGCGGU	3617
345	TGACCGCA	GGCTAGCTACAACGA	GTGTTAGG	2908	CCUAACAC	A UGCGGUCA	3618
347	AGTGACCG	GGCTAGCTACAACGA	ATGTTGTA	2909	UAACACAU	G CGGUCACU	3619
350	AAAAGTGA	GGCTAGCTACAACGA	CGCATGTG	2910	CACAUGCG	G UCACUUUU	3620
353	CCCAAAAG	GGCTAGCTACAACGA	GACCGCAT	2911	AUGCGGUC	A CUUUUGGG	3621
364	CCACACCA	GGCTAGCTACAACGA	CTCCCAAA	2912	UUUGGGAG	A UGGUGUGG	3622
367	CTCCACAA	GGCTAGCTACAACGA	CATCTCCC	2913	GGGAGAUG	G UGUGGGAG	3623
369	TGCTTCCA	GGCTAGCTACAACGA	ACCATCTC	2914	GAGAUGGU	G UGGGAGCA	3624
375	CTTTTCTG	GGCTAGCTACAACGA	TCCACAC	2915	GUGUGGGA	G CAGAAAAG	3625
383	CACCCCTG	GGCTAGCTACAACGA	TTTTCTGC	2916	GCAGAAAA	G CAGGGGUG	3626
389	TGACGACA	GGCTAGCTACAACGA	CCCTGCTT	2917	AAGCAGGG	G UGUUGUCA	3627
391	CATGACGA	GGCTAGCTACAACGA	ACCCCTGC	2918	GCAGGGGU	G UCGUCAUG	3628
394	GAGCATGA	GGCTAGCTACAACGA	GACACCCC	2919	GGGGUGUC	G UCAUGCUC	3629
397	GTTGAGCA	GGCTAGCTACAACGA	GACGACAC	2920	GUGUCGUC	A UGCUCAAC	3630
399	CTGTTGAG	GGCTAGCTACAACGA	ATGACGAC	2921	GUCGUCAU	G CUCAACAG	3631
404	TCACTCTG	GGCTAGCTACAACGA	TGAGCATG	2922	CAUGCUCU	A CAGAGUGA	3632
409	CTCCATCA	GGCTAGCTACAACGA	TCTGTTGA	2923	UCAACAGA	G UGAUGGAG	3633
412	TTTCTCCA	GGCTAGCTACAACGA	CACCTGTT	2924	ACAGAGUG	A UGGAGAAA	3634

Table 6

422	TTAACGAA	GGCTAGCTACACGA	CTTTCTCC	2925	GGAGAAAG G	UUCGUUAA	3635
426	CATTTTAA	GGCTAGCTACACGA	GAACCTTT	2926	AAAGGUUC G	UUA AAAAUG	3636
432	TGTGCGCA	GGCTAGCTACACGA	TTTAACGA	2927	UCGUUAAA A	UGCGCACA	3637
434	ATTGTGCG	GGCTAGCTACACGA	ATTTTAAC	2928	GUUAAA AU	G CGCACA AU	3638
436	GTATTGTG	GGCTAGCTACACGA	GCATTTTA	2929	UAAA AUGC G	CACAAUAC	3639
438	CAGTATTG	GGCTAGCTACACGA	GCGCATTT	2930	AA AUGCGC A	CAAUACUG	3640
441	GCCAGTA	GGCTAGCTACACGA	TGTGCGCA	2931	UGCGCACA A	UACUGGCC	3641
443	GTGGCCAG	GGCTAGCTACACGA	ATTGTGCG	2932	CGCACAAU A	CUGGCCAC	3642
447	TTTTGTGG	GGCTAGCTACACGA	CAGTATTG	2933	CAAUACUG G	CCACAAAA	3643
450	TCTTTTTG	GGCTAGCTACACGA	GGCCAGTA	2934	UACUGGCC A	CAAAAAGA	3644
469	AAAGATCA	GGCTAGCTACACGA	CTCTTTTT	2935	AAAAGAG A	UGAUCUUU	3645
472	TTCAAAGA	GGCTAGCTACACGA	CATCTCTT	2936	AAAGAU G A	UCUUUGAA	3646
482	AATTGTG	GGCTAGCTACACGA	CTTCAAAG	2937	CUUUGAAG A	CACAAAUU	3647
484	CAAAATTTG	GGCTAGCTACACGA	GTCTTCAA	2938	UUGAAGAC A	CAAAUUUG	3648
488	ATTTCAAA	GGCTAGCTACACGA	TTGTGTCT	2939	AGACACAA A	UUUGAAAU	3649
495	AATGTTAA	GGCTAGCTACACGA	TTCAAAAT	2940	AAUUUGAA A	UUAACA AU	3650
499	GATCAATG	GGCTAGCTACACGA	TAATTTCA	2941	UGAAA UUA A	CAUUGAUC	3651
501	GAGATCAA	GGCTAGCTACACGA	GTTAATTT	2942	AAAUUAAC A	UUGAUCUC	3652
505	TTCAGAGA	GGCTAGCTACACGA	CAATGTTA	2943	UAACAUTG A	UCUCUGAA	3653
515	ACTTGATA	GGCTAGCTACACGA	CTTCAGAG	2944	CUCUGAAG A	UAUCAAGU	3654
517	TGACTTGA	GGCTAGCTACACGA	ATCTTCAG	2945	CUGAAGAU A	UCAAGUCA	3655
522	TAATATGA	GGCTAGCTACACGA	TTGATATC	2946	GAUAUCAA G	UCAUAUUA	3656
525	GTATAATA	GGCTAGCTACACGA	GACTTGAT	2947	AUCAAGUC A	UAUUUAUC	3657
527	CTGTATAA	GGCTAGCTACACGA	ATGACTTG	2948	CAAGUCAU A	UUAUACAG	3658
530	GCACTGTA	GGCTAGCTACACGA	AATATGAC	2949	GUCAUAUU A	UACAGUGC	3659
532	TGCACTG	GGCTAGCTACACGA	ATAATATG	2950	CAUAUAU A	CAGUGCGA	3660
535	CTGTGCGA	GGCTAGCTACACGA	TGTATAAT	2951	AUAUAACA G	UGCGACAG	3661
537	AGCTGTG	GGCTAGCTACACGA	ACTGTATA	2952	UAUACAGU G	CGACAGCU	3662
540	TCTAGCTG	GGCTAGCTACACGA	CGCACTGT	2953	ACAGUGCG A	CAGCUAGA	3663
543	AATTCTAG	GGCTAGCTACACGA	TGTCGCAC	2954	GUGCGACA G	CUAGAAUU	3664
549	TTTTCCAA	GGCTAGCTACACGA	TCTAGCTG	2955	CAGCUAGA A	UUGGAAAA	3665

Table 6

557	TTGTAAGG	GGCTAGCTACAACGA	TTTCCAAT	2956	AUUGGAAA	A	CCUUAACA	3666
562	TTGGGTTG	GGCTAGCTACAACGA	AGGTTTTT	2957	AAAACCUU	A	CAACCCAA	3667
565	TTCTTGGG	GGCTAGCTACAACGA	TGTAAGGT	2958	ACCUUACA	A	CCCAAGAA	3668
574	CTCTCGAG	GGCTAGCTACAACGA	TTCTTGGG	2959	CCCAAGAA	A	CUCGAGAG	3669
583	ATGTAAGA	GGCTAGCTACAACGA	CTCTCGAG	2960	CUCGAGAG	A	UCUUAACU	3670
588	TGGAAATG	GGCTAGCTACAACGA	AAGATCTC	2961	GAGAUUUU	A	CAUUUCCA	3671
590	AGTGGAAA	GGCTAGCTACAACGA	GTAAGATC	2962	GAUCUUAC	A	UUUCCACU	3672
596	TGGTATAG	GGCTAGCTACAACGA	GGAAATGT	2963	ACAUUCC	A	CUAUACCA	3673
599	ATGTGGTA	GGCTAGCTACAACGA	AGTGGAAA	2964	UUUCCACU	A	UACCACAU	3674
601	CCATGTGG	GGCTAGCTACAACGA	ATAGTGGA	2965	UCCACUAA	A	CCACAUGG	3675
604	AGGCCATG	GGCTAGCTACAACGA	GGTATAGT	2966	ACUUAACC	A	CAUGGCCU	3676
606	TCAGGCCA	GGCTAGCTACAACGA	GTGGTATA	2967	UAUACCAC	A	UGGCCUGA	3677
609	AAGTCAGG	GGCTAGCTACAACGA	CATGTGGT	2968	ACCACAUG	G	CCUGACUU	3678
614	CTCCAAAG	GGCTAGCTACAACGA	CAGGCCAT	2969	AUGGCCUG	A	CUUUGGAG	3679
622	TTCAGGGA	GGCTAGCTACAACGA	TCCAAAGT	2970	ACUUUGGA	G	UCCCUGAA	3680
630	GCTGGTGA	GGCTAGCTACAACGA	TCAGGGAC	2971	GUCCUGA	A	UCACCAGC	3681
633	GAGGCTGG	GGCTAGCTACAACGA	GATTCAGG	2972	CCUGAAUC	A	CCAGCCUC	3682
637	GAATGAGG	GGCTAGCTACAACGA	TGGTGATT	2973	AAUACCCA	G	CCUCAUUC	3683
642	TTCNAGAA	GGCTAGCTACAACGA	GAGGCTGG	2974	CCAGCCUC	A	UUCUUGAA	3684
650	AAAGAAAG	GGCTAGCTACAACGA	TCAAGAAT	2975	AUUCUUUGA	A	CUUUCUUU	3685
664	CTCTCGGA	GGCTAGCTACAACGA	TTTGAANA	2976	UUUUCAAA	G	UCCGAGAG	3686
672	GACCCCTGA	GGCTAGCTACAACGA	TCTCGGAC	2977	GUCCGAGA	G	UCAGGGUC	3687
678	CTGAGTGA	GGCTAGCTACAACGA	CCTGACTC	2978	GAGUCAGG	G	UCACUCAG	3688
681	GGGCTGAG	GGCTAGCTACAACGA	GACCCTGA	2979	UCAGGGUC	A	CUCAGCCC	3689
686	GCTCCGGG	GGCTAGCTACAACGA	TGAGTGAC	2980	GUCACUCA	G	CCCGGAGC	3690
693	GGCCCGTG	GGCTAGCTACAACGA	TCCGGGCT	2981	AGCCCGGA	G	CACGGGCC	3691
695	CGGGCCCC	GGCTAGCTACAACGA	GCTCCGGG	2982	CCCGGAGC	A	CGGGCCCC	3692
699	ACAACGGG	GGCTAGCTACAACGA	CCGTGCTC	2983	GAGCACGG	G	CCCGUUGU	3693
703	CACCACAA	GGCTAGCTACAACGA	GGGCCCGT	2984	ACGGGCCC	G	UUGUGGUG	3694
706	GTGCACCA	GGCTAGCTACAACGA	AACGGGCC	2985	GGCCCGUU	G	UGGUGCAC	3695
709	GCAGTGCA	GGCTAGCTACAACGA	CACAACGG	2986	CCGUUGUG	G	UGCACUGC	3696

Table 6

711	CTGCAGTG	GGCTAGCTACAACGA	ACCACAAC	2987	GUUGUGGU G	CACUGCAG	3697
713	CACTGCAG	GGCTAGCTACAACGA	GCACCACA	2988	UGUGGUGC A	CUGCAGUG	3698
716	CTGCACTG	GGCTAGCTACAACGA	AGTGCACC	2989	GGUGCACU G	CAGUGCAG	3699
719	TGCCTGCA	GGCTAGCTACAACGA	TGCAGTGC	2990	GCACUGCA G	UGCAGGCA	3700
721	GATGCCCTG	GGCTAGCTACAACGA	ACTGCAGT	2991	ACUGCAGU G	CAGGCAUC	3701
725	TGCCGATG	GGCTAGCTACAACGA	CTGCAC TG	2992	CAGUGCAG G	CAUCGGCA	3702
727	CCTGCCGA	GGCTAGCTACAACGA	GCCTGCAC	2993	GUGCAGGC A	UCGCAGG	3703
731	CAGACCTG	GGCTAGCTACAACGA	CGATGCCT	2994	AGGCAUCG G	CAGGUCUG	3704
735	GTTCCAGA	GGCTAGCTACAACGA	CTGCCGAT	2995	AUCGGCAG G	UCUGGAAC	3705
742	ACAGAAGG	GGCTAGCTACAACGA	TCCAGACC	2996	GGUCUGGA A	CCUUCUGU	3706
749	CAGCCAGA	GGCTAGCTACAACGA	AGAAGGTT	2997	AACCUUCU G	UCUGGCUG	3707
754	GGTATCAG	GGCTAGCTACAACGA	CAGACAGA	2998	UCUGUCUG G	CUGAUACC	3708
758	GGCAGGTA	GGCTAGCTACAACGA	CAGCCAGA	2999	UCUGGCUG A	UACCGGCC	3709
760	GAGGCAGG	GGCTAGCTACAACGA	ATCAGCCA	3000	UGGCUGAU A	CCUGCCUC	3710
764	GCAAGAGG	GGCTAGCTACAACGA	AGGTATCA	3001	UGAUACCU G	CCUCUUGC	3711
771	TCCATCAG	GGCTAGCTACAACGA	AAGAGGCA	3002	UGCCUCUU G	CUGAUGGA	3712
775	CTTGTCCA	GGCTAGCTACAACGA	CAGCAAGA	3003	UCUUGCUG A	UGGACAAG	3713
779	TCCTCTTG	GGCTAGCTACAACGA	CCATCAGC	3004	GCUGAUGG A	CAAGAGGA	3714
791	AAGAAGGG	GGCTAGCTACAACGA	CTTTCCTC	3005	GAGGAAAG A	CCCUCUCU	3715
802	GATATCAA	GGCTAGCTACAACGA	GGAGGAG	3006	CUUCUUCG G	UUGAUUUC	3716
806	TCTTGATA	GGCTAGCTACAACGA	CAACGGNA	3007	UUCCGUUG A	UAUCAAGA	3717
808	TTTCTTGA	GGCTAGCTACAACGA	ATCAACGG	3008	CCGUUGAU A	UCRAGAAA	3718
817	TAACAGCA	GGCTAGCTACAACGA	TTTCTTGA	3009	UCAAGAAA G	UGCUGUUA	3719
819	TCTAACAG	GGCTAGCTACAACGA	ACTTCTTT	3010	AAGAAAGU G	CUGUUAGA	3720
822	ATTCTTAA	GGCTAGCTACAACGA	AGCACTTT	3011	AAAGUGCU G	UUAGAAAU	3721
829	CTTCTTCA	GGCTAGCTACAACGA	TTCTAACA	3012	UGUUAGAA A	UGAGGAAG	3722
837	ATCCGAAA	GGCTAGCTACAACGA	TTCTCTAT	3013	AUGAGGAA G	UUUCGGAU	3723
844	CAGCCCCA	GGCTAGCTACAACGA	CGAAACT	3014	AGUUUCGG A	UGGGGCUG	3724
849	TGGATCAG	GGCTAGCTACAACGA	CCCATCCG	3015	CGGAUGGG G	CUGAUCCA	3725
853	TGCTTGGA	GGCTAGCTACAACGA	CAGCCCCA	3016	UGGGGCUG A	UCCAGACA	3726
859	GTCGGCTG	GGCTAGCTACAACGA	CTGGATCA	3017	UGAUCCAG A	CAGCCGAC	3727

Table 6

862	CTGGTCGG	GGCTAGCTACAACGA	TGCTCTGA	3018	UCCAGACA G CCGACCAG	3728
866	GCAGCTGG	GGCTAGCTACAACGA	CGGCTGTC	3019	GACAGCCG A CCAGCUGC	3729
870	AAGCGCAG	GGCTAGCTACAACGA	TGGTCGGC	3020	GCCGACCA G CUGCGCUU	3730
873	GAGAAGCG	GGCTAGCTACAACGA	AGCTGGTC	3021	GACCAGCU G CGUUCUC	3731
875	AGGAGAAG	GGCTAGCTACAACGA	GCAGCTGG	3022	CCAGCUGC G CUUCUCCU	3732
884	CAGCCAGG	GGCTAGCTACAACGA	AGGAGAAG	3023	CUUCUCCU A CCUGGCUG	3733
889	GATCACAG	GGCTAGCTACAACGA	CAGGTAGG	3024	CCUACCUG G CUGUGAUC	3734
892	TTCGATCA	GGCTAGCTACAACGA	AGCCAGGT	3025	ACCUGGCU G UGAUCGAA	3735
895	ACCTTCGA	GGCTAGCTACAACGA	CACAGCCA	3026	UGGCUGUG A UCGAAGGU	3736
902	ATTTGGCA	GGCTAGCTACAACGA	CTTCGATC	3027	GAUGAAG G UGCCAAAU	3737
904	GAATTTGG	GGCTAGCTACAACGA	ACCTTCGA	3028	UCGAAGGU G CCAAUUC	3738
909	ATGATGAA	GGCTAGCTACAACGA	TTGGCACC	3029	GGUGCCAA A UUCAUCAU	3739
913	CCCCATGA	GGCTAGCTACAACGA	GAATTTGG	3030	CCAAUUC A UCAUGGGG	3740
916	GTCCCCCA	GGCTAGCTACAACGA	GATGAATT	3031	AAUUCAU A UGGGGGAC	3741
923	CGGAAGAG	GGCTAGCTACAACGA	CCCCCATG	3032	CAUGGGG A CUCUUCGG	3742
931	ATCCTGCA	GGCTAGCTACAACGA	GGAAGAGT	3033	ACUCUUC G UGCAGGAU	3743
933	TGATCCTG	GGCTAGCTACAACGA	ACGGAAGA	3034	UCUUCGU G CAGGAUCA	3744
938	TCCACTGA	GGCTAGCTACAACGA	CCTGCACG	3035	CGUGCAGG A UCAGUGGA	3745
942	TCCTTTCA	GGCTAGCTACAACGA	TGATCCTG	3036	CAGGAUCA G UGGAAGGA	3746
951	TGGGAAAG	GGCTAGCTACAACGA	TCCTTTCA	3037	UGGAAGGA G CUUCCCCA	3747
959	GGTCCTCG	GGCTAGCTACAACGA	GGGAAGC	3038	GUUUCUCC A CGAGGACC	3748
965	GCTCCAGG	GGCTAGCTACAACGA	CCTCGTGG	3039	CCAGGAGG A CCUGGAGC	3749
972	GGTGGGGG	GGCTAGCTACAACGA	TCCAGTTC	3040	GACCUGGA G CCCCACC	3750
978	TGCTCGGG	GGCTAGCTACAACGA	GGGGGCTC	3041	GAGCCCCC A CCCGAGCA	3751
984	GGGATATG	GGCTAGCTACAACGA	TCGGGTGG	3042	CCACCCTGA G CAUAUCCC	3752
986	GGGGGATA	GGCTAGCTACAACGA	GCTCGGCT	3043	ACCCGAGC A UAUCCCCC	3753
988	TGGGGGGA	GGCTAGCTACAACGA	ATGCTCGG	3044	CCGAGCAU A UCCCCCCA	3754
996	CGGGGAGG	GGCTAGCTACAACGA	GGGGGAT	3045	AUCCCCC A CCUCCCCG	3755
1005	TTGGGTGG	GGCTAGCTACAACGA	CGGGGAGG	3046	CCUCCCCG G CCACCCAA	3756
1008	CGTTTGGG	GGCTAGCTACAACGA	GGCCGGGG	3047	CCCGGCC A CCCAAACG	3757
1014	AGGATTCG	GGCTAGCTACAACGA	TTGGGTGG	3048	CCACCCAA A CGAAUCCU	3758

Table 6

1018	CTCCAGGA	GGCTAGCTACAACGA	TCGTTTGG	3049	CCAAACGA A UCCUGGAG	3759
1026	TTGTGTGG	GGCTAGCTACAACGA	TCCAGGAT	3050	AUCCUGGA G CCACACAA	3760
1029	CCATTGTG	GGCTAGCTACAACGA	GGCTCCAG	3051	CUGGAGCC A CACAAUGG	3761
1031	TCCCATTTG	GGCTAGCTACAACGA	GTGCTCC	3052	GGAGCCAC A CAAUGGGA	3762
1034	ATTTCCCA	GGCTAGCTACAACGA	TGTGTGGC	3053	GCCACACA A UGGGAAAU	3763
1041	TCCCTGCA	GGCTAGCTACAACGA	TTCCCATTT	3054	AAUGGGAA A UGCAGGGA	3764
1043	ACTCCCTG	GGCTAGCTACAACGA	ATTTCCCA	3055	UGGGAUUU G CAGGGAGU	3765
1050	GGGAAGAA	GGCTAGCTACAACGA	TCCCTGCA	3056	UGCAGGGA G UUCUCCCC	3766
1061	ACTGGTGA	GGCTAGCTACAACGA	TTGGGAAG	3057	CUUCCCAA A UCACCAGU	3767
1064	CCCCTGG	GGCTAGCTACAACGA	GATTTGGG	3058	CCCAAUUC A CCAGUGGG	3768
1068	TTACACCA	GGCTAGCTACAACGA	TGTTGATT	3059	AAUCACCA G UGGGUGAA	3769
1072	TTCTTTCA	GGCTAGCTACAACGA	CCACTGGT	3060	ACCAGUGG G UGAAGGAA	3770
1084	CTCCTGGG	GGCTAGCTACAACGA	CTCTTCCT	3061	AGGAAGAG A CCCAGGAG	3771
1094	AGTCTTTA	GGCTAGCTACAACGA	CCTCTGG	3062	CCAGGAGG A UAAAGACU	3772
1100	TGGGCAG	GGCTAGCTACAACGA	CTTTATCC	3063	GGUAAAAG A CUGCCCCA	3773
1103	TGATGGGG	GGCTAGCTACAACGA	AGTCTTTA	3064	UAAAGACU G CCCCACUA	3774
1108	TTCTTTGA	GGCTAGCTACAACGA	GGGCGAGT	3065	ACUGCCCC A UCAAGGAA	3775
1127	TTAAGGGG	GGCTAGCTACAACGA	TTCTTTTT	3066	AAAAGGAA G CCCCUUAA	3776
1136	GTGCGCA	GGCTAGCTACAACGA	TTAAGGGG	3067	CCCCUUA A UGCCGCAC	3777
1138	GGGTGCGG	GGCTAGCTACAACGA	ATTTAAGG	3068	CCUUAUU G CCGCACCC	3778
1141	GTAGGGTG	GGCTAGCTACAACGA	GGCATTTA	3069	UAAUUGCC G CACCCUAC	3779
1143	CGTAGGGG	GGCTAGCTACAACGA	GCGGCATT	3070	AAUGCCGC A CCUACGG	3780
1148	CGATGCCG	GGCTAGCTACAACGA	AGGTTGCG	3071	CGCACCCU A CCGCAUCG	3781
1151	TTTCGATG	GGCTAGCTACAACGA	CGTAGGGT	3072	ACCCUACG G CAUCGAAA	3782
1153	GCTTTTCA	GGCTAGCTACAACGA	GCCGTAGG	3073	CCUACGSC A UCGAAAGC	3783
1160	GACTCATG	GGCTAGCTACAACGA	TTTCGATG	3074	CAUCGAAA G CAUGAGUC	3784
1162	TTGACTCA	GGCTAGCTACAACGA	GCTTTTCA	3075	UCGAAAGC A UGAGUCA	3785
1166	TGCTTTGA	GGCTAGCTACAACGA	TCATGCTT	3076	AAGCAUGA G UCAAGACA	3786
1172	CTTCAGTG	GGCTAGCTACAACGA	CTTGACTC	3077	GAGUCAAG A CACUGAAG	3787
1174	AACCTCAG	GGCTAGCTACAACGA	GTCCTTAC	3078	GUCNAGAC A CUGAAGUU	3788
1180	ACTTCTAA	GGCTAGCTACAACGA	TTCAGTGT	3079	ACACUGAA G UAGAAGU	3789

Table 6

1187	CGACCCGA	GGCTAGCTACAACGA	TTCTAACT	3080	AGUAGAA G UCGGUGG	3790
1192	CCCCACGA	GGCTAGCTACAACGA	CCGACTTC	3081	GAAGUCGG G UCGUGGGG	3791
1195	TCCCCCCA	GGCTAGCTACAACGA	GACCCGAC	3082	GUCGGGUC G UGGGGGGA	3792
1205	CTCGAAGA	GGCTAGCTACAACGA	TTCCCCCC	3083	GGGGGGA G UCUUCGAG	3793
1214	CCTGGGCA	GGCTAGCTACAACGA	CTCGAAGA	3084	UCUUCGAG G UGCCCAGG	3794
1216	AGCCTGGG	GGCTAGCTACAACGA	ACCTCGAA	3085	UUCGAGGU G CCCAGGCU	3795
1222	GGAGGCAG	GGCTAGCTACAACGA	CTGGGCAC	3086	GUGCCCG G CUGCCUCC	3796
1225	TGGGGAGG	GGCTAGCTACAACGA	AGCCTGGG	3087	CCAGGGCU G CCUCCCA	3797
1234	CCCTTTGG	GGCTAGCTACAACGA	TGGGGAGG	3088	CCUCCCA G CCAAAGGG	3798
1245	AGTGACGG	GGCTAGCTACAACGA	TCCCTTTT	3089	AAAGGGGA G CCGUCACU	3799
1248	GGCAGTGA	GGCTAGCTACAACGA	GGCTCCCC	3090	GGGGAGCC G UCACUGCC	3800
1251	TCGGGCAG	GGCTAGCTACAACGA	GACGGCTC	3091	GAGCCGUC A CUGCCCGA	3801
1254	TTCTCGGG	GGCTAGCTACAACGA	AGTGACGG	3092	CCGUCACU G CCCGAGAA	3802
1265	GGTCCTCG	GGCTAGCTACAACGA	CCTTCTCG	3093	CGAGNAGG A CGAGGACC	3803
1271	GTGCATGG	GGCTAGCTACAACGA	CCTCGTCC	3094	GGAGGAGG A CCAUGCAC	3804
1274	TCAGTGCA	GGCTAGCTACAACGA	GGTCCTCG	3095	CGAGGACC A UGCACUGA	3805
1276	ACTCAGTG	GGCTAGCTACAACGA	ATGGTCCT	3096	AGGACCAU G CACUGAGU	3806
1278	TAACTCAG	GGCTAGCTACAACGA	GCATGGTC	3097	GACCAUGC A CUGAGUUA	3807
1283	TCCAGTAA	GGCTAGCTACAACGA	TCAGTGCA	3098	UGCACUGA G UUACUGGA	3808
1286	GCTTCCAG	GGCTAGCTACAACGA	AACTCAGT	3099	ACUGAGUT A CUGGAAGC	3809
1293	AGGAAGGG	GGCTAGCTACAACGA	TTCCAGTA	3100	UACUGGAA G CCCUUCCT	3810
1303	CATGTTGA	GGCTAGCTACAACGA	CAGGAAGG	3101	CCUCCUG G UCAACAUG	3811
1307	CGCACATG	GGCTAGCTACAACGA	TAACCAGG	3102	CCUGGUCA A CAUGUGCG	3812
1309	CACGCACA	GGCTAGCTACAACGA	GTTGACCA	3103	UGGUCAAC A UGUGCGUG	3813
1311	GCCACGCA	GGCTAGCTACAACGA	ATGTTGAC	3104	GUCAACAU G UGCGUGGC	3814
1313	TAGCCACG	GGCTAGCTACAACGA	ACATGTTG	3105	CAACAUGU G CGUGGCUA	3815
1315	CGTAGCCA	GGCTAGCTACAACGA	GCACATGT	3106	ACAUGUGC G UGGCUACG	3816
1318	GACCGTAG	GGCTAGCTACAACGA	CAGGCACA	3107	UGUGCGUG G CUACGGUC	3817
1321	GAGGACCG	GGCTAGCTACAACGA	AGCCACGC	3108	GCGUGGCU A CGGUCCUC	3818
1324	CGTGAGGA	GGCTAGCTACAACGA	CGTAGCCA	3109	UGGCUACG G UCCUCACG	3819
1330	GCCGGCCG	GGCTAGCTACAACGA	GAGGACCG	3110	CGGUCCUC A CGGCCGCG	3820

Table 6

1333	AGCGCCGG	GGCTAGCTACAACGA	CGTGAGGA	3111	UCCUACG G CCGCGCU	3821
1337	GGTAAGCG	GGCTAGCTACAACGA	CGGCCGTG	3112	CACGCCG G CGUUACC	3822
1339	GAGGTAAG	GGCTAGCTACAACGA	GCCGGCCG	3113	CGGCCGG G CUUACCUC	3823
1343	AGCAGAGG	GGCTAGCTACAACGA	AGCGCCG	3114	CGCGCUU A CCUCUGCU	3824
1349	ACCTGTAG	GGCTAGCTACAACGA	AGAGGTAA	3115	UUACCUCU G CUACAGGU	3825
1352	GGAACCTG	GGCTAGCTACAACGA	AGCAGAGG	3116	CCUCUGCU A CAGGUUCC	3826
1356	AACAGGAA	GGCTAGCTACAACGA	CTGTAGCA	3117	UGCUACAG G UUCCUGUU	3827
1362	CTGTTGAA	GGCTAGCTACAACGA	AGGAACCT	3118	AGGUUCCU G UUCAACAG	3828
1367	TGTTGCTG	GGCTAGCTACAACGA	TGAACAGG	3119	CCUGUUA A CAGCAACA	3829
1370	ATGTGTTG	GGCTAGCTACAACGA	TGTTGAAC	3120	GUUCACA G CAACACAU	3830
1373	GCTATGTG	GGCTAGCTACAACGA	TGCTGTTG	3121	CAACAGCA A CACAUAGC	3831
1375	AGGCTATG	GGCTAGCTACAACGA	GTGCTGT	3122	ACAGCAAC A CAUAGCCU	3832
1377	TCAGGCTA	GGCTAGCTACAACGA	GTGTTGCT	3123	AGCAACAC A UAGCCUGA	3833
1380	GGGTCAGG	GGCTAGCTACAACGA	TATGTGTT	3124	AACACAUA G CCUGACCC	3834
1385	GAGGAGGG	GGCTAGCTACAACGA	CAGGCTAT	3125	AUAGCCUG A CCUCCUC	3835
1395	AGGTGGAG	GGCTAGCTACAACGA	GGAGGAGG	3126	CCUCCUCC A CUCCACCU	3836
1400	GGTGGAGG	GGCTAGCTACAACGA	GGAGTGGA	3127	UCCACUCC A CCUCCACC	3837
1406	ACAGTGGG	GGCTAGCTACAACGA	GGAGGTGG	3128	CCACUCC A CCCACUGU	3838
1410	GCGGACAG	GGCTAGCTACAACGA	GGGTGGAG	3129	CUCCACC A CUGUCCGC	3839
1413	GAGCGGA	GGCTAGCTACAACGA	AGTGGGTG	3130	CACCCACU G UCCGCCUC	3840
1417	GGCAGAGG	GGCTAGCTACAACGA	GGACAGTG	3131	CACUGUCC G CCUCUGCC	3841
1423	TCTGCGGG	GGCTAGCTACAACGA	AGAGGCGG	3132	CCGCCUCU G CCGCAGA	3842
1427	GGGCTCTG	GGCTAGCTACAACGA	GGGCAGAG	3133	CUCUGCCG G CAGAGCCC	3843
1432	GGCGTGGG	GGCTAGCTACAACGA	TCTGCGGG	3134	CCCGCAGA G CCACACGC	3844
1436	GTCGGGCG	GGCTAGCTACAACGA	GGGCTCTG	3135	CAGAGCCC A CGCCCCGAC	3845
1438	TAGTCGGG	GGCTAGCTACAACGA	GTGGGCTC	3136	GAGCCAC G CCGACUA	3846
1443	CCTGTAG	GGCTAGCTACAACGA	CGGGCGTG	3137	CAGCCCG A CUAGCAGG	3847
1447	CATGCCTG	GGCTAGCTACAACGA	TAGTCGGG	3138	CCGACUA G CAGGCAUG	3848
1451	GCGGCATG	GGCTAGCTACAACGA	CTGTAGT	3139	ACUAGCAG G CAUGCCGC	3849
1453	CCGGGGCA	GGCTAGCTACAACGA	GCCTGCTA	3140	UAGCAGGC A UGCGCGG	3850
1455	TACCGCGG	GGCTAGCTACAACGA	ATGCTGTC	3141	GCAGGCAU G CCGCGGUA	3851

Table 6

1458	ACCTACCG	GGCTAGCTACAACGA	GGCATGCC	3142	GGCAUGCC G	CGGUAGGU	3852
1461	CTTACCTA	GGCTAGCTACAACGA	CGCGGCAT	3143	AUGCCGCG G	UAGGUAAG	3853
1465	GGCCCTTA	GGCTAGCTACAACGA	CTACCGCG	3144	CGCGGUAG G	UAAGGGCC	3854
1471	TCCGGCGG	GGCTAGCTACAACGA	CCTTACCT	3145	AGGUAAGG G	CCGCCGGA	3855
1474	CGGTCCGG	GGCTAGCTACAACGA	GGCCCTTA	3146	UAAAGGCC G	CCGGACCG	3856
1479	CTACGCGG	GGCTAGCTACAACGA	CGCGCGC	3147	GCCGCCG A	CCGCGUAG	3857
1482	TCTCTACG	GGCTAGCTACAACGA	GGTCCGCG	3148	GCCGGACC G	CGUAGAGA	3858
1484	GCTCTCTA	GGCTAGCTACAACGA	GCGTCCG	3149	CGGACCGC G	UAGAGAGC	3859
1491	GGGCCCGG	GGCTAGCTACAACGA	TCTCTACG	3150	CGUAGAGA G	CCGGGCC	3860
1496	GTCCGGGG	GGCTAGCTACAACGA	CCGGCTCT	3151	AGAGCCGG G	CCCCGGAC	3861
1503	AACGTCCG	GGCTAGCTACAACGA	CGGGGCC	3152	GGCCCCGG A	CGGACGUT	3862
1507	AACCAACG	GGCTAGCTACAACGA	CGTCCGG	3153	CCGGACGG A	CGUUGGUT	3863
1509	AGAACCAA	GGCTAGCTACAACGA	GTCCGTCC	3154	GGAGGCAC G	UUGGUUCU	3864
1513	GTGCAGAA	GGCTAGCTACAACGA	CAACGTCC	3155	GGAGGUUG G	UUCUGCAC	3865
1518	TTTTAGTG	GGCTAGCTACAACGA	AGAACCAA	3156	UUGGUUCU G	CACUAAAA	3866
1520	GGTTTTAG	GGCTAGCTACAACGA	GCAGAAC	3157	GGUUCUGC A	CCCAUCUU	3867
1526	AAGATGGG	GGCTAGCTACAACGA	TTTAGTGC	3158	GCACUAAA A	CCCAUCUU	3868
1530	GGGAAGA	GGCTAGCTACAACGA	GGTTTITA	3159	UAAAACCC A	UCUUCGCC	3869
1541	GACACACA	GGCTAGCTACAACGA	CCGGGGA	3160	UUCGCCGG A	UGUGUGUC	3870
1543	GAGACACA	GGCTAGCTACAACGA	ATCCGGGG	3161	CCCGGAU G	UGUGUCUC	3871
1545	GTGAGACA	GGCTAGCTACAACGA	ACATCCGG	3162	CCGGAUGU G	UCUCACCC	3872
1547	GGGTGAGA	GGCTAGCTACAACGA	ACACATCC	3163	GGAGUGU G	UCUCACCC	3873
1552	ATGAGGGG	GGCTAGCTACAACGA	GAGACACA	3164	UGUGUCUC A	CCCCUCAU	3874
1559	TAAAAGGA	GGCTAGCTACAACGA	GAGGGGTG	3165	CACCCUC A	UCCUUUUA	3875
1567	GCAAAAAG	GGCTAGCTACAACGA	AAAAGGAT	3166	AUCCUUU A	CUUUUUGC	3876
1574	GGAAGGGG	GGCTAGCTACAACGA	AAAAGTA	3167	UACUUUUU G	CCCCUUC	3877
1583	ACTCAAAG	GGCTAGCTACAACGA	GGAAGGGG	3168	CCCCUCC A	CUUUGAGU	3878
1590	ATTTGGTA	GGCTAGCTACAACGA	TCAAAGTG	3169	CACUUUGA G	UACCAAAU	3879
1592	GGATTTGG	GGCTAGCTACAACGA	ACTCAAAG	3170	CUUUGAGU A	CCAAAUCC	3880
1597	CTTGTGGA	GGCTAGCTACAACGA	TTGGTACT	3171	AGUACCAA A	UCCACAAG	3881
1601	ATGGCTTG	GGCTAGCTACAACGA	GGATTTGG	3172	CCAAAUCC A	CAAGCCAU	3882

Table 6

1605	AAAAATGG	GGCTAGCTACAACGA	TTGTGGAT	3173	AUCCACAA G CCAUUUUU	3883
1608	TCAAAAAA	GGCTAGCTACAACGA	GGCTTGTG	3174	CACAAGCC A UUUUUUGA	3884
1622	CTCTTTCA	GGCTAGCTACAACGA	TCTCTCTCA	3175	UGAGGAGA G UGAAAGAG	3885
1632	GCATGGTA	GGCTAGCTACAACGA	TCTCTTTC	3176	GAAAGAGA G UACCAUGC	3886
1634	CAGCATGG	GGCTAGCTACAACGA	ACTCTCTT	3177	AAGAGAGU A CCAUGCUG	3887
1637	CGCCAGCA	GGCTAGCTACAACGA	GGTACTCT	3178	AGAGUACC A UGCUGGCG	3888
1639	GCCGCCAG	GGCTAGCTACAACGA	ATGGTACT	3179	AGUACCAU G CUGGCGGC	3889
1643	CTGCGCCG	GGCTAGCTACAACGA	CAGCATGG	3180	CCAUGCUG G CGGCGCAG	3890
1646	CCTCTGCG	GGCTAGCTACAACGA	CGCCAGCA	3181	UGCUGGCG G CGCAGAGG	3891
1648	TCCCTCTG	GGCTAGCTACAACGA	GCCGCCAG	3182	CUGGCGGC G CAGAGGGA	3892
1661	GGTGTAGG	GGCTAGCTACAACGA	CCCTTCCC	3183	GGGAAAGG G CCUACACC	3893
1665	GACGGGTG	GGCTAGCTACAACGA	AGGCCCTT	3184	AGGGGCCU A CACCCGUC	3894
1667	AAGACGGG	GGCTAGCTACAACGA	GTAGGCCC	3185	GGGCCUAC A CCCGUCUU	3895
1671	CCCCAAGA	GGCTAGCTACAACGA	GGGTGTAG	3186	CUACACCC G UCUGGGG	3896
1679	GGGGCGAG	GGCTAGCTACAACGA	CCCAAGAC	3187	GUCUUGG G CUCGCCCC	3897
1683	GGGTGGGG	GGCTAGCTACAACGA	GAGCCCCA	3188	UGGGGCUC G CCCCACCC	3898
1688	GCCCTGGG	GGCTAGCTACAACGA	GGGGCGAG	3189	CUCGCCCC A CCCAGGGC	3899
1695	GGAGGGAG	GGCTAGCTACAACGA	CCTGGGTG	3190	CACCCAGG G CUCCCUCC	3900
1708	CTGGGATG	GGCTAGCTACAACGA	TCCAGGAG	3191	CUCCUGGA G CAUCCCAG	3901
1710	GCCTGGGA	GGCTAGCTACAACGA	GCTCCAGG	3192	CCUGGAGC A UCCCAGGC	3902
1717	GCCGCCCG	GGCTAGCTACAACGA	CTGGGATG	3193	CAUCCAG G CGGGCGGC	3903
1721	GCGTGCCG	GGCTAGCTACAACGA	CCGCTGTG	3194	CCAGGGG G CGGCACGC	3904
1724	CTGGCGTG	GGCTAGCTACAACGA	CGCCCGCC	3195	GGCGGGCG G CACGCCAG	3905
1726	GTCTGGCG	GGCTAGCTACAACGA	GCCGCCCG	3196	CGGGCGGC A CGCCAGAC	3906
1728	CTGTCTGG	GGCTAGCTACAACGA	GTGCGGCC	3197	GGCGGCAC G CCAGACAG	3907
1733	GGGGCTTG	GGCTAGCTACAACGA	CTGGCGTG	3198	CACGCCAG A CAGCCCCC	3908
1736	GGGGGGGG	GGCTAGCTACAACGA	TGTCTGGC	3199	GCCAGACA G CCCCCCCC	3909
1749	CCTGCAGA	GGCTAGCTACAACGA	TCAAGGGG	3200	CCCCUUGA A UCUGCAGG	3910
1753	GCTCCCTG	GGCTAGCTACAACGA	AGATTCAA	3201	UUGAAUCU G CAGGGAGC	3911
1760	GAGAGTTG	GGCTAGCTACAACGA	TCCCTGCA	3202	UGCAGGGA G CAACUCUC	3912
1763	GTGGAGAG	GGCTAGCTACAACGA	TGCTCCCT	3203	AGGGAGCA A CUCUCCAC	3913

Table 6

1770	ATATGGAG	GGCTAGCTACAACGA	GGAGAGTT	3204	AACUCUCC A CUCCAUAU	3914
1775	AATAAATA	GGCTAGCTACAACGA	GGAGTGGA	3205	UCCACUCC A UAUUUUUU	3915
1777	TAAATAAA	GGCTAGCTACAACGA	ATGGAGTG	3206	CACUCCAU A UUUUUUUA	3916
1781	TGTTTAAA	GGCTAGCTACAACGA	AAATATGG	3207	CCAUAUUU A UUUAAAACA	3917
1787	AAAAATTG	GGCTAGCTACAACGA	TTAATAAA	3208	UUUUUUAA A CAAUUUUU	3918
1790	GGAAAAAA	GGCTAGCTACAACGA	TGTTTAAA	3209	UUUAAACA A UUUUUUCC	3919
1805	TATGGATG	GGCTAGCTACAACGA	CTTTGGGG	3210	CCCCAAAG G CAUCCAUA	3920
1807	ACTATGGA	GGCTAGCTACAACGA	GCCTTTGG	3211	CCAAAGGC A UCCAUAUG	3921
1811	GTGCACTA	GGCTAGCTACAACGA	GGATGCCT	3212	AGGCAUCC A UAGUGCAC	3922
1814	CTAGTGCA	GGCTAGCTACAACGA	TATGGATG	3213	CAUCCAUA G UGCACUAG	3923
1816	TGCTAGTG	GGCTAGCTACAACGA	ACTATGGA	3214	UCCAUAUG G CACUAGCA	3924
1818	AATGCTAG	GGCTAGCTACAACGA	GCACTATG	3215	CAUAGUGC A CUAGCAUU	3925
1822	AGAAATG	GGCTAGCTACAACGA	TAGTGCAC	3216	GUGCACUA G CAUUUUUU	3926
1824	CAAGAAAA	GGCTAGCTACAACGA	GCTAGTGC	3217	GCACUAGC A UUUUUUUG	3927
1834	ATTATTGG	GGCTAGCTACAACGA	TCAAGAAA	3218	UUUUUUGA A CCAAUAAU	3928
1838	ATACATTA	GGCTAGCTACAACGA	TGGTTCAA	3219	UUGAACCA A UAAUGUAU	3929
1841	TTAATACA	GGCTAGCTACAACGA	TATTGGTT	3220	AACCAAUA A UGUUUUAA	3930
1843	TTTAAATA	GGCTAGCTACAACGA	ATTATTGG	3221	CCAUAUUU G UAUUAAAA	3931
1845	AATTTTAA	GGCTAGCTACAACGA	ACATTATT	3222	AAUAAUGU A UUAAAAUU	3932
1851	TCAAAAAA	GGCTAGCTACAACGA	TTAATATC	3223	GUUUUAAA A UUUUUUGA	3933
1859	GGCTGACA	GGCTAGCTACAACGA	CAAAAAAT	3224	AUUUUUUG A UGUCAGCC	3934
1861	AAGGCTGA	GGCTAGCTACAACGA	ATCAAAAA	3225	UUUUUUGA G UCAGCCUU	3935
1865	ATGCAAGG	GGCTAGCTACAACGA	TGACATCA	3226	UGAUGUCA G CCUUGCAU	3936
1870	CCTTGATG	GGCTAGCTACAACGA	AAGGCTGA	3227	UCAGCCUU G CAUCAAGG	3937
1872	GCCCTTGA	GGCTAGCTACAACGA	GCAAGGCT	3228	AGCCUUGC A UCAAGGGC	3938
1879	TGATAAAG	GGCTAGCTACAACGA	CCTTGATG	3229	CAUCAAGG G CUUUUAUA	3939
1884	CTTTTGTG	GGCTAGCTACAACGA	AAAGCCCT	3230	AGGCGUUU A UCAAAAAAG	3940
1892	TTATTGTA	GGCTAGCTACAACGA	TTTTTGAT	3231	AUCAAAAA G UACAAUAA	3941
1894	TATTATTG	GGCTAGCTACAACGA	ACTTTTGG	3232	CAAAAAGU A CAAUAAUA	3942
1897	ATTATTAT	GGCTAGCTACAACGA	TGTACTTT	3233	AAAGUACA A UAAUAAAU	3943
1900	AGGATTTA	GGCTAGCTACAACGA	TATTGTAC	3234	GUACAAUA A UAAAUCCU	3944

Table 6

1904	CCTGAGGA	GGCTAGCTACAACGA	TTATTATT	3235	AAUAAUAA A UCCUCAGG	3945
1912	CAGTACTA	GGCTAGCTACAACGA	CTGAGGAT	3236	AUCCUCAG G UAGUACUG	3946
1915	TCCCAGTA	GGCTAGCTACAACGA	TACCTGAG	3237	CUCAGGUA G UACUGGGA	3947
1917	ATTCCCAG	GGCTAGCTACAACGA	ACTACCTG	3238	CAGGUAGU A CUGGGAUU	3948
1924	GCCTTCCA	GGCTAGCTACAACGA	TCCCAGTA	3239	UACUGGGA A UGGAAGGC	3949
1931	TGGCAAAG	GGCTAGCTACAACGA	CTTCCATT	3240	AAUGGAAG G CUUUGCCA	3950
1936	GCCCATGG	GGCTAGCTACAACGA	AAAGCCTT	3241	AAGGCUUU G CCAUGGGC	3951
1939	CAGGCCCA	GGCTAGCTACAACGA	GGCAAAGC	3242	GCUUUGCC A UGGGCCUG	3952
1943	GCAGCAGG	GGCTAGCTACAACGA	CCATGGCA	3243	UGCCAUGG G CCUGCUGC	3953
1947	TGACGCAG	GGCTAGCTACAACGA	AGGCCCAT	3244	AUGGGCCU G CUGCGUCA	3954
1950	GTCTGACG	GGCTAGCTACAACGA	AGCAGGCC	3245	GGCCUGCU G CGUCAGAC	3955
1952	TGGTCTGA	GGCTAGCTACAACGA	GCAGCAGG	3246	CCUGCUGC G UCAGACCA	3956
1957	AGTACTGG	GGCTAGCTACAACGA	CTGACGCA	3247	UGCGUCAG A CCAGUACU	3957
1961	TCCCAGTA	GGCTAGCTACAACGA	TGGTCTGA	3248	UCAGACCA G UACUGGGA	3958
1963	CTTCCCAG	GGCTAGCTACAACGA	ACTGGTCT	3249	AGACCAGU A CUGGGAAG	3959
1976	TACAACCG	GGCTAGCTACAACGA	CCTCCTTC	3250	GAAGGAGG A CGUUGUA	3960
1979	GCTTACAA	GGCTAGCTACAACGA	CGTCTCTC	3251	GGAGGACG G UUGUAAGC	3961
1982	ACTGCTTA	GGCTAGCTACAACGA	AACCGTCC	3252	GGACGGUU G UAAGCAGU	3962
1986	AACAAC TG	GGCTAGCTACAACGA	TTACAACC	3253	GGUUGUAA G CAGUUGUU	3963
1989	AATAACAA	GGCTAGCTACAACGA	TGCTTACA	3254	UGUAAAGCA G UUGUUUAU	3964
1992	CTAAATAA	GGCTAGCTACAACGA	AACTGCTT	3255	AAGCAGUU G UUAUUUAG	3965
1995	TCACTAAA	GGCTAGCTACAACGA	AACAAC TG	3256	CAGUUGUU A UUUAGUGA	3966
2000	CAATATCA	GGCTAGCTACAACGA	TAAATAAC	3257	GUUAUUUA G UGAUAUUG	3967
2003	CCACAATA	GGCTAGCTACAACGA	CACTAAAT	3258	AUUUAGUG A UAUUGUGG	3968
2005	ACCCACAA	GGCTAGCTACAACGA	ATCACTAA	3259	UUAGUGAU A UUGUGGGU	3969
2008	GTTACCCA	GGCTAGCTACAACGA	AATATCAC	3260	GUGAUUUU G UGGGUUAC	3970
2012	TCACGTTA	GGCTAGCTACAACGA	CCACAATA	3261	UAUUGUGG G UAACGUGA	3971
2015	TTCTCACG	GGCTAGCTACAACGA	TACCCACA	3262	UGUGGGUA A CGUGAGAA	3972
2017	TCTTCTCA	GGCTAGCTACAACGA	GTTACCCA	3263	UGGGUAAAC G UGAGAAGA	3973
2025	TGTTTCTA	GGCTAGCTACAACGA	CTTCTCAC	3264	GUGAGAAG A UAGAACAA	3974
2030	TAGCATTG	GGCTAGCTACAACGA	TCTATCTT	3265	AAGAUAGA A CAAUGCUA	3975

Table 6

2033	TTATAGCA	GGCTAGCTACAACGA	TGTTCTAT	3266	AUAGAACA	A UGUUAUA	3976
2035	TATTATAG	GGCTAGCTACAACGA	ATTGTTCT	3267	AGAACA	AU G CUUAUA	3977
2038	ATATATTA	GGCTAGCTACAACGA	AGCATTGT	3268	ACAAUGCU	A UAAUAUAU	3978
2041	ATTATATA	GGCTAGCTACAACGA	TATAGCAT	3269	AUGCUAUA	A UAUUAUAU	3979
2043	TCATTATA	GGCTAGCTACAACGA	ATTATAGC	3270	GUUAUAU	A UAUUAUA	3980
2045	GTTTATT	GGCTAGCTACAACGA	ATATTATA	3271	UAUAUAU	A UAAUAAC	3981
2048	CGTGTTC	GGCTAGCTACAACGA	TATATATT	3272	AAUAUAU	A UGAACACG	3982
2052	CCACAGTG	GGCTAGCTACAACGA	TCATTATA	3273	UAUAUAU	A CACGUGG	3983
2054	TACCCACG	GGCTAGCTACAACGA	GTTTATT	3274	UAUAUAU	A CGUGGUA	3984
2056	AATACCCA	GGCTAGCTACAACGA	GTGTTTAT	3275	AUGAACAC	G UGGUUAU	3985
2060	ATTAAATA	GGCTAGCTACAACGA	CCACGTGT	3276	ACACGUGG	G UAUUAUAU	3986
2062	TTATTAAA	GGCTAGCTACAACGA	ACCCACGT	3277	ACGUGGGU	A UUAUAUA	3987
2067	GTTTCTTA	GGCTAGCTACAACGA	TAAATACC	3278	GGUAUUUA	A UAAGAAAC	3988
2074	ACATCATG	GGCTAGCTACAACGA	TTCTTATT	3279	AAUAAGAA	A CAUGAUGU	3989
2076	TCACATCA	GGCTAGCTACAACGA	GTTTCTTA	3280	UAAGAAAC	A UGAUGUA	3990
2079	ATCTCACA	GGCTAGCTACAACGA	CATGTTTC	3281	GAACAUG	A UGUGAGU	3991
2081	TAATCTCA	GGCTAGCTACAACGA	ATCATGTT	3282	AAUAUAU	G UGAGAUUA	3992
2086	CAAAGTAA	GGCTAGCTACAACGA	CTCACATC	3283	GAUGUGAG	A UUAUUUG	3993
2089	GGACAAAG	GGCTAGCTACAACGA	AATCTCAC	3284	GUGAGAUU	A CUUUGUCC	3994
2094	AAGGGGA	GGCTAGCTACAACGA	AAAGTAAT	3285	AUAACUUU	G UCCCGCUU	3995
2099	AGAATAAG	GGCTAGCTACAACGA	GGACAAA	3286	UUUGUCCC	G CUUAUUU	3996
2103	GAGCAGAA	GGCTAGCTACAACGA	AAGGGGA	3287	UCCCGCUU	A UUCUGCUC	3997
2108	ACAGGGAG	GGCTAGCTACAACGA	AGAATAAG	3288	CUUAUUU	G CUCCCGU	3998
2115	GCAGATAA	GGCTAGCTACAACGA	AGGAGCA	3289	UGCUCCU	G UUAUCUGC	3999
2118	CTAGCAGA	GGCTAGCTACAACGA	AACAGGGA	3290	UCCUGUU	A UCUGCUAG	4000
2122	AGATCTAG	GGCTAGCTACAACGA	AGATAACA	3291	UGUAUUU	G CUAGAUCU	4001
2127	GAAC TAGA	GGCTAGCTACAACGA	CTAGCAGA	3292	UCUGCUAG	A UCUAGUUC	4002
2132	ATTGAGAA	GGCTAGCTACAACGA	TAGATCTA	3293	UAGAUCUA	G UUCUCAU	4003
2139	AGCAGTGA	GGCTAGCTACAACGA	TGAGAACT	3294	AGUUCUCA	A UCACUGCU	4004
2142	GGAGCAG	GGCTAGCTACAACGA	GATTGAGA	3295	UCUCAUC	A CUGCUCC	4005
2145	CGGGGAG	GGCTAGCTACAACGA	AGTGATTG	3296	CAUAUCU	G CUCCCCG	4006

Table 6

2153	AATACACA	GGCTAGCTACAACGA	GGGGGAGC	3297	GUUUUUUU G UGUUUUUU	4007
2155	CTAATACA	GGCTAGCTACAACGA	ACGGGGGA	3298	UUUUUUUU G UGUUUUUU	4008
2157	TTCTAATA	GGCTAGCTACAACGA	ACACGGGG	3299	UUUUUUUU G UGUUUUUU	4009
2159	CATTCTAA	GGCTAGCTACAACGA	ACACACGG	3300	UUUUUUUU G UGUUUUUU	4010
2165	TACATGCA	GGCTAGCTACAACGA	TCTAATAC	3301	UUUUUUUU G UGUUUUUU	4011
2167	CTTACATG	GGCTAGCTACAACGA	ATTCTAAT	3302	UUUUUUUU G UGUUUUUU	4012
2169	ACCTTACA	GGCTAGCTACAACGA	GCATTCTA	3303	UUUUUUUU G UGUUUUUU	4013
2171	AGACCTTA	GGCTAGCTACAACGA	ATGCATTCT	3304	UUUUUUUU G UGUUUUUU	4014
2176	CAAGAAGA	GGCTAGCTACAACGA	CTTACATG	3305	UUUUUUUU G UGUUUUUU	4015
2184	TCAGGACA	GGCTAGCTACAACGA	AAGAAGAC	3306	UUUUUUUU G UGUUUUUU	4016
2186	CATCAGGA	GGCTAGCTACAACGA	ACAAGAAG	3307	UUUUUUUU G UGUUUUUU	4017
2192	ATTTTTC	GGCTAGCTACAACGA	CAGGACAC	3308	UUUUUUUU G UGUUUUUU	4018
2199	AGCACATA	GGCTAGCTACAACGA	TTTTCATC	3309	UUUUUUUU G UGUUUUUU	4019
2201	CAAGCACA	GGCTAGCTACAACGA	ATTTTTC	3310	UUUUUUUU G UGUUUUUU	4020
2203	TTCAAGCA	GGCTAGCTACAACGA	ATTTTTC	3311	UUUUUUUU G UGUUUUUU	4021
2205	ATTTCAAG	GGCTAGCTACAACGA	ACATATTT	3312	UUUUUUUU G UGUUUUUU	4022
2212	GTTTCTCA	GGCTAGCTACAACGA	TTTTCATC	3313	UUUUUUUU G UGUUUUUU	4023
2219	GATCAAG	GGCTAGCTACAACGA	TTTTCATC	3314	UUUUUUUU G UGUUUUUU	4024
2225	AGCAGAGA	GGCTAGCTACAACGA	CAAAGTTT	3315	UUUUUUUU G UGUUUUUU	4025
2231	TTAGTAAG	GGCTAGCTACAACGA	AGAGATCA	3316	UUUUUUUU G UGUUUUUU	4026
2235	CACATTAG	GGCTAGCTACAACGA	AAGCAGAG	3317	UUUUUUUU G UGUUUUUU	4027
2239	GGGGCACA	GGCTAGCTACAACGA	TAGTAAGC	3318	UUUUUUUU G UGUUUUUU	4028
2241	ATGGGGCA	GGCTAGCTACAACGA	ATTAGTAA	3319	UUUUUUUU G UGUUUUUU	4029
2243	ACATGGGG	GGCTAGCTACAACGA	ACATTAGT	3320	UUUUUUUU G UGUUUUUU	4030
2248	CTTGGACA	GGCTAGCTACAACGA	GGGGCACA	3321	UUUUUUUU G UGUUUUUU	4031
2250	GACTTTGA	GGCTAGCTACAACGA	ATGGGGCA	3322	UUUUUUUU G UGUUUUUU	4032
2256	AGTTTGA	GGCTAGCTACAACGA	TTGGACAT	3323	UUUUUUUU G UGUUUUUU	4033
2261	CAGGCAGG	GGCTAGCTACAACGA	TGGACTTG	3324	UUUUUUUU G UGUUUUUU	4034
2265	TGCACAGG	GGCTAGCTACAACGA	AGTTTGA	3325	UUUUUUUU G UGUUUUUU	4035
2269	GTCATGCA	GGCTAGCTACAACGA	AGGCAGGT	3326	UUUUUUUU G UGUUUUUU	4036
2271	AGGTCATG	GGCTAGCTACAACGA	ACAGGCAG	3327	UUUUUUUU G UGUUUUUU	4037

Table 6

2273	TCAGGTCA	GGCTAGCTACAACGA	GCACAGGC	3328	GCCUGUC A	UGACCUGA	4038
2276	TGATCAGG	GGCTAGCTACAACGA	CATGCACA	3329	UGUGCAUG A	CCUGAUC A	4039
2281	TGTAATGA	GGCTAGCTACAACGA	CAGGTCAT	3330	AUGACCUG A	UCAUUACA	4040
2284	CCATGTAA	GGCTAGCTACAACGA	GATCAGGT	3331	ACCUGAUC A	UUACAUGG	4041
2287	CAGCCATG	GGCTAGCTACAACGA	AATGATCA	3332	UGAUAUU A	CAUGGCUG	4042
2289	CACAGCCA	GGCTAGCTACAACGA	GTAATGAT	3333	AUCAUAC A	UGGCUGUG	4043
2292	AACCACAG	GGCTAGCTACAACGA	CATGTAAT	3334	AUUACAUG G	CUGUGGUU	4044
2295	AGGAACCA	GGCTAGCTACAACGA	AGCCATGT	3335	ACAUGGCU G	UGGUUCCU	4045
2298	CTTAGGAA	GGCTAGCTACAACGA	CACAGCCA	3336	UGGCUGUG G	UUCCUAAG	4046
2306	GCAACAGG	GGCTAGCTACAACGA	TTAGGAAC	3337	GUUCCUAA G	CCUGUUGC	4047
2310	TTACAGCA	GGCTAGCTACAACGA	AGGCTTAG	3338	CUAAGCCU G	UUGCUGAA	4048
2313	GACTTCAG	GGCTAGCTACAACGA	AACAGGCT	3339	AGCCUGUU G	CUGAAGUC	4049
2319	GACAAATGA	GGCTAGCTACAACGA	TTACAGAA	3340	UUGCUGAA G	UCAUUGUC	4050
2322	AGCGACAA	GGCTAGCTACAACGA	GACTTCAG	3341	CUGAAGUC A	UUGUCGCU	4051
2325	CTGAGCGA	GGCTAGCTACAACGA	AATGACTT	3342	AAGUCAUU G	UGGCUCAG	4052
2328	TTGCTGAG	GGCTAGCTACAACGA	GACAAATGA	3343	UCAUUGUC G	CUCAGCAA	4053
2333	CCCTATTG	GGCTAGCTACAACGA	TGAGCGAC	3344	GUCGCUCA G	CAAUAGGG	4054
2336	GCACCCCTA	GGCTAGCTACAACGA	TGCTGAGC	3345	GCUCAGCA A	UAGGGUUC	4055
2341	AAACTGCA	GGCTAGCTACAACGA	CCTATTGC	3346	GCAUAGG G	UGCAGUUU	4056
2343	GAAAACTG	GGCTAGCTACAACGA	ACCTATT	3347	AAUAGGGU G	CAGUUUUC	4057
2346	CTGGAAAA	GGCTAGCTACAACGA	TGCACCCCT	3348	AGGUGCA G	UUUCCAG	4058
2357	AATGCCTA	GGCTAGCTACAACGA	TCTGGAA	3349	UUCAGGA A	UAGGCAUU	4059
2361	GGCAAAATG	GGCTAGCTACAACGA	CTATTCTT	3350	AGGAAUAG G	CAUUGGCC	4060
2363	TAGGCAAA	GGCTAGCTACAACGA	GCCTATT	3351	GAAUAGGC A	UUUGCCUA	4061
2367	GAATTAGG	GGCTAGCTACAACGA	AAATGCCT	3352	AGGCAUUU G	CCUAAUUC	4062
2372	GCCAGGAA	GGCTAGCTACAACGA	TAGGCAAA	3353	UUUGCCUA A	UUCCUGGC	4063
2379	GTGTCATG	GGCTAGCTACAACGA	CAGGAATT	3354	AAUCCUG G	CAUGACAC	4064
2381	GAGTGTC A	GGCTAGCTACAACGA	GCCAGGAA	3355	UUCCUGGC A	UGACACUC	4065
2384	CTAGAGTG	GGCTAGCTACAACGA	CATGCCAG	3356	CUGGCAUG A	CACUCUAG	4066
2386	CACTAGAG	GGCTAGCTACAACGA	GTCATGCC	3357	GGCAUGAC A	CUCUAGUG	4067
2392	GGAAGTCA	GGCTAGCTACAACGA	TAGAGTGT	3358	ACACUCUA G	UGACUUCC	4068

Table 6

2395	CCAGGAAG	GGCTAGCTACAACGA	CACTAGAG	3359	CUCUAGUG	A CUCUCCUGG	4069
2403	GGGCCTCA	GGCTAGCTACAACGA	CAGGAAGT	3360	ACUUCUG	G UGAGCCCC	4070
2408	AGGCTGGG	GGCTAGCTACAACGA	CTCACCAG	3361	CUGGUGAG	G CCCAGCCU	4071
2413	AGGACAGG	GGCTAGCTACAACGA	TGGCCTC	3362	GAGGCCCA	G CCUGUCCU	4072
2417	TACCAGGA	GGCTAGCTACAACGA	AGGCTGGG	3363	CCCAGCCU	G UCCUGGUA	4073
2423	CTGCTGTA	GGCTAGCTACAACGA	CAGGACAG	3364	CUGUCCUG	G UACAGCAG	4074
2425	CCCTGCTG	GGCTAGCTACAACGA	ACCAGGAC	3365	GUCCUGGU	A CAGCAGGG	4075
2428	AGACCCCTG	GGCTAGCTACAACGA	TGTACCAG	3366	CUGGUACA	G CAGGGUCU	4076
2433	CAGCAAGA	GGCTAGCTACAACGA	CCTGCTGT	3367	ACAGCAGG	G UCUUGCUG	4077
2438	AGTTACAG	GGCTAGCTACAACGA	AAGACCTT	3368	AGGGUCUU	G CUGUAACU	4078
2441	CTGAGTTA	GGCTAGCTACAACGA	AGCAAGAC	3369	GUCUUGCU	G UAACUCAG	4079
2444	TGTCTGAG	GGCTAGCTACAACGA	TACAGCAA	3370	UUGCUGUA	A CUCAGACA	4080
2450	TTGGAATG	GGCTAGCTACAACGA	CTGAGTTA	3371	UAACUCAG	A CAUCCAA	4081
2452	CCTTGGAA	GGCTAGCTACAACGA	GTCTGAGT	3372	ACUCAGAC	A UUCCAAGG	4082
2461	TTCCCATTA	GGCTAGCTACAACGA	CCTTGGAA	3373	UUCCAAGG	G UAUGGAA	4083
2463	GCTTCCCA	GGCTAGCTACAACGA	ACCTTGGG	3374	CCAAGGGU	A UGGGAAGC	4084
2470	GAATATGG	GGCTAGCTACAACGA	TTCCCATTA	3375	UAUGGGAA	G CCAUAUUC	4085
2473	TGTGAATA	GGCTAGCTACAACGA	GGCTTCCC	3376	GGGAAGCC	A UAUUCACA	4086
2475	GGTGTGAA	GGCTAGCTACAACGA	ATGGCTTC	3377	GAAGCCAU	A UUCACACC	4087
2479	GTGAGGTG	GGCTAGCTACAACGA	GAATATGG	3378	CCAUAUUC	A CACCUCAC	4088
2481	GCGTGAGG	GGCTAGCTACAACGA	GTGAATAT	3379	AUAUUCAC	A CCUCACGC	4089
2486	CCAGAGCG	GGCTAGCTACAACGA	GAGGTGTG	3380	CACACCUC	A CGCUCUGG	4090
2488	GTCCAGAG	GGCTAGCTACAACGA	GTGAGGTG	3381	CACCUCAC	G CUCUGGAC	4091
2495	AAATCATG	GGCTAGCTACAACGA	CCAGAGCG	3382	CGCUCUGG	A CAUGAUUU	4092
2497	CTAAATCA	GGCTAGCTACAACGA	GTCCAGAG	3383	CUCUGGAC	A UGAUUUAG	4093
2500	TCCCTAAA	GGCTAGCTACAACGA	CATGTCCA	3384	UGGACAUG	A UUUAGGGA	4094
2510	TGTCCCTG	GGCTAGCTACAACGA	TTCCCTAA	3385	UUAGGGAA	G CAGGGACA	4095
2516	GGGGGGTG	GGCTAGCTACAACGA	CCCTGTCT	3386	AAGCAGGG	A CACCCCCC	4096
2518	GCGGGGGG	GGCTAGCTACAACGA	GTCCCTGC	3387	GCAGGGAC	A CCCCCCGC	4097
2525	GTGGGGGG	GGCTAGCTACAACGA	GGGGGGTG	3388	CACCCCCC	G CCCCCCAC	4098
2532	CCCAAAGG	GGCTAGCTACAACGA	GGGGGGGG	3389	CGCCCCCC	A CCUUUGGG	4099

Table 6

2541	GAGGCTGA	GGCTAGCTACAACGA	CCCAAAGG	3390	CCUUUGG A UCAGCCUC	4100
2545	GGCGGAGG	GGCTAGCTACAACGA	TGATCCCA	3391	UGGGAUCA G CCUCCGCC	4101
2551	TGGAATGG	GGCTAGCTACAACGA	GGAGGCTG	3392	CAGCCUCC G CCAUCCCA	4102
2554	ACTTGGAA	GGCTAGCTACAACGA	GGGGGAGG	3393	CCUCCGCC A UUCCAAGU	4103
2561	AGTGTCTGA	GGCTAGCTACAACGA	TTGGAATG	3394	CAUCCAA G UCGACACU	4104
2565	GAAGAGTG	GGCTAGCTACAACGA	CGACTTGG	3395	CCAAGUCG A CACUCUUC	4105
2567	AAGAAGAG	GGCTAGCTACAACGA	GTCGACTT	3396	AAGUGAC A CUCUUCUU	4106
2578	ACGCTCTG	GGCTAGCTACAACGA	TCAAGAAG	3397	CUUCUUGA G CAGACCGU	4107
2582	AATCACGG	GGCTAGCTACAACGA	CTGCTCAA	3398	UUGAGCAG A CCGUGAUU	4108
2585	CCAAATCA	GGCTAGCTACAACGA	GGTCTGCT	3399	AGCAGACC G UGAUUUGG	4109
2588	CTTCCAAA	GGCTAGCTACAACGA	CACGGTCT	3400	AGACCGUG A UUUGGAAG	4110
2601	AGCAGGTG	GGCTAGCTACAACGA	CTCTCTTC	3401	GAAGAGAG G CACCUGCU	4111
2603	CCAGCAGG	GGCTAGCTACAACGA	GCCTCTCT	3402	AGAGAGGC A CCUGCUGG	4112
2607	GTTTCCAG	GGCTAGCTACAACGA	AGGTGCCT	3403	AGGCACCU G CUGGAAAC	4113
2614	AAGTGTGG	GGCTAGCTACAACGA	TTCAGACA	3404	UGCUGGAA A CCACACUU	4114
2617	AAGAAGTG	GGCTAGCTACAACGA	GGTTTCCA	3405	UGGAAACC A CACUUCUU	4115
2619	TCAAGAAG	GGCTAGCTACAACGA	GTGGTTTC	3406	GAACCCAC A CUUCUUGA	4116
2629	CCAGGCTG	GGCTAGCTACAACGA	TTCAAGAA	3407	UUCUUGAA A CAGCCUGG	4117
2632	CACCCAGG	GGCTAGCTACAACGA	TGTTTCAA	3408	UUGAAACA G CCUGGGUG	4118
2638	GACCGTCA	GGCTAGCTACAACGA	CCAGGCTG	3409	CAGCCUGG G UGACGGUC	4119
2641	AAGACCG	GGCTAGCTACAACGA	CACCCAGG	3410	CCUGGGUG A CGGUCCUU	4120
2644	CTAAAGGA	GGCTAGCTACAACGA	CGTCACCC	3411	GGGUGAGC G UCCUUUAG	4121
2653	GCAGGCTG	GGCTAGCTACAACGA	CTAAAGGA	3412	UCCUUAG G CAGCCUGC	4122
2656	GCGGCAGG	GGCTAGCTACAACGA	TGCTTAAA	3413	UUUAGGCA G CCUGCCGC	4123
2660	GACGCGG	GGCTAGCTACAACGA	AGGCTGCC	3414	GGCAGCCU G CCGCCGUC	4124
2663	AGAGACGG	GGCTAGCTACAACGA	GGCAGGCT	3415	AGCCUGCC G CCGUCUCU	4125
2666	GACAGAGA	GGCTAGCTACAACGA	GGCGGCAG	3416	CUGCCGCC G UCUCUGUC	4126
2672	AACCGGGA	GGCTAGCTACAACGA	AGAGACGG	3417	CCGUCUCU G UCCCGGUU	4127
2678	AAGGTGAA	GGCTAGCTACAACGA	CGGGACAG	3418	CUGUCCCG G UUCACCUU	4128
2682	CGGCAAGG	GGCTAGCTACAACGA	GAACGGGG	3419	CCCGGUUC A CCUUGCCG	4129
2687	TCTCTCGG	GGCTAGCTACAACGA	AAGGTGAA	3420	UUCACCUU G CCGAGAGA	4130

Table 6

2697	CAGACGG GGCTAGCTACAACGA CTCTCTCG	3421	CGAGAGAG G CGGUCUG	4131
2699	GGCAGAG GGCTAGCTACAACGA GCCTCTCT	3422	AGAGAGGC G CGUCUGCC	4132
2701	GGGGCAGA GGCTAGCTACAACGA GCGCTCT	3423	AGAGGCGC G UCUGCCCC	4133
2705	GGGTGGGG GGCTAGCTACAACGA AGACGGCC	3424	GCGGUCU G CCCACACC	4134
2710	TTTGAGGG GGCTAGCTACAACGA GGGGCAGA	3425	UCUGCCCC A CCCUCAA	4135
2718	CCACAGGG GGCTAGCTACAACGA TTGAGGGT	3426	ACCCUCAA A CCCUGUGG	4136
2723	AGCCCCCA GGCTAGCTACAACGA AGGGTTTG	3427	CAACCCU G UGGGCCU	4137
2728	CCATCAGG GGCTAGCTACAACGA CCCACAGG	3428	CCUGUGG G CCUGAUGG	4138
2733	GAGCACCA GGCTAGCTACAACGA CAGGCCCC	3429	GGGGCCUG A UGGUGCUC	4139
2736	CGTGAGCA GGCTAGCTACAACGA CATCAGGC	3430	GCCUGAUG G UGCUCACG	4140
2738	GTCGTGAG GGCTAGCTACAACGA ACCATCAG	3431	CUGAUGG G CUCACGAC	4141
2742	AAGAGTCG GGCTAGCTACAACGA GAGCACCA	3432	UGGUGCUC A CGACUCUU	4142
2745	AGGAAGAG GGCTAGCTACAACGA CGTGAGCA	3433	UGCUCACG A CUCUUCUU	4143
2754	TCCCTTTG GGCTAGCTACAACGA AGGAAGAG	3434	CUCUUCU G CAAAGGGA	4144
2763	GTCCTTCA GGCTAGCTACAACGA TCCCTTTG	3435	CAAAGGGA A CUGAAGAC	4145
2770	TGTGGAGG GGCTAGCTACAACGA CTTGAGTT	3436	AACUGAAG A CCUCCACA	4146
2776	ACTTAATG GGCTAGCTACAACGA GGAGGTCT	3437	AGACCUCC A CAUUAAGU	4147
2778	CCACTTAA GGCTAGCTACAACGA GTGGAGGT	3438	ACCUCCAC A UUAAGUGG	4148
2783	AAAAGCCA GGCTAGCTACAACGA TTAATGTG	3439	CACAUUAA G UGGCUUUU	4149
2786	TTAAAAG GGCTAGCTACAACGA CACTTAAT	3440	AUUAAGUG G CUUUUUA	4150
2794	TTTTCATG GGCTAGCTACAACGA TAAAAGC	3441	GCUUUUUA A CAUGAAAA	4151
2796	GTTTTTCA GGCTAGCTACAACGA GTTAAAA	3442	UUUUUAAC A UGAAAAAC	4152
2803	CTGCCGTG GGCTAGCTACAACGA TTTTCATG	3443	CAUGAAAA A CACGGCAG	4153
2805	AGTGCCG GGCTAGCTACAACGA GTTTTTCA	3444	UGAAAAAC A CGGCAGCU	4154
2808	TACAGCTG GGCTAGCTACAACGA CGTGTGTT	3445	AAAACAG G CAGCUGUA	4155
2811	AGCTACAG GGCTAGCTACAACGA TGCCGTGT	3446	ACACGGCA G CUGUAGCU	4156
2814	GGGAGCTA GGCTAGCTACAACGA AGCTGCCG	3447	CGGCAGCU G UAGCUCCC	4157
2817	CTCGGGAG GGCTAGCTACAACGA TACAGCTG	3448	CAGCUGUA G CUCCCCAG	4158
2825	GAGAGTAG GGCTAGCTACAACGA TCGGGAGC	3449	GCUCGCCA G CUACUCUC	4159
2828	CAAGAGAG GGCTAGCTACAACGA AGTCTGGG	3450	CCCGAGCU A CUCUCUUG	4160
2836	AATGCTGG GGCTAGCTACAACGA AAGAGAGT	3451	ACUCUCUU G CCAGCAUU	4161

Table 6

2840	TGAAAATG	GGCTAGCTACACGA	TGGCAAGA	3452	UCUUGCCA G CAUUUUA	4162
2842	TGTGAAA	GGCTAGCTACACGA	GCTGGCAA	3453	UUGCAGC A UUUUCACA	4163
2848	GCAAAATG	GGCTAGCTACACGA	GAAAATGC	3454	GCAUUUC A CAUUUUGC	4164
2850	AGGCAAAA	GGCTAGCTACACGA	GTGAAAAT	3455	AUUUCAC A UUUUGCCU	4165
2855	GAGAAAGG	GGCTAGCTACACGA	AAAATGTG	3456	CACAUUU G CCUUUCUC	4166
2864	TTCTACCA	GGCTAGCTACACGA	GAGAAAGG	3457	CCUUUCUC G UGGUAGAA	4167
2867	GGCTTCTA	GGCTAGCTACACGA	CACGAGAA	3458	UUCUCGUG G UAGAAGCC	4168
2873	TGTACTGG	GGCTAGCTACACGA	TTCTACCA	3459	UGGUAGAA G CCAGUACA	4169
2877	TCTCTGTA	GGCTAGCTACACGA	TGGCTTCT	3460	AGAAAGCA G UACAGAGA	4170
2879	TTTCTCTG	GGCTAGCTACACGA	ACTGGCTT	3461	AAGCCAGU A CAGAGAAA	4171
2887	CCACAGAA	GGCTAGCTACACGA	TTCTCTGT	3462	ACAGAGAA A UUCUGUGG	4172
2892	TCCCACCA	GGCTAGCTACACGA	AGAAATTC	3463	GAAAUUCU G UGGUGGGA	4173
2895	TGTTCCCA	GGCTAGCTACACGA	CACAGAAT	3464	AUUCUGUG G UGGGAACA	4174
2901	CTCGAATG	GGCTAGCTACACGA	TCCCACCA	3465	UGGUGGGA A CAUUCGAG	4175
2903	ACCTCGAA	GGCTAGCTACACGA	GTTCCAC	3466	GUGGGAAC A UUCGAGGU	4176
2910	GGGTGACA	GGCTAGCTACACGA	CTCGAATG	3467	CAUUCGAG G UGUCACCC	4177
2912	CAGGTGTA	GGCTAGCTACACGA	ACCTCGAA	3468	UUCGAGGU G UCACCCUG	4178
2915	CTGCAGGG	GGCTAGCTACACGA	GACACCTC	3469	GAGGUGUC A CCCUGCAG	4179
2920	TAGCTCTG	GGCTAGCTACACGA	AGGCTGAC	3470	GUCACCCU G CAGAGCUA	4180
2925	CACCATAG	GGCTAGCTACACGA	TTGTCAGG	3471	CCUGCAGA G CUAUGGUG	4181
2928	CCTCACCA	GGCTAGCTACACGA	AGCTCTGC	3472	GCAGAGCU A UGGUGAGG	4182
2931	ACACCTCA	GGCTAGCTACACGA	CATAGCTC	3473	GAGCUAUG G UGAGGUGU	4183
2936	TATCCACA	GGCTAGCTACACGA	CTCACCAT	3474	AUGGUGAG G UGUGGAUA	4184
2938	CTTATCCA	GGCTAGCTACACGA	ACCTCACC	3475	GGUGAGGU G UGGAUAG	4185
2942	AAGCCTTA	GGCTAGCTACACGA	CCACACCT	3476	AGGUGUGG A UAAGGCUU	4186
2947	CACCTAAG	GGCTAGCTACACGA	CTTATCCA	3477	UGGAUAAG G CUUAGGUG	4187
2953	GCCTGGCA	GGCTAGCTACACGA	CTAAGCCT	3478	AGGCUUAG G UGCCAGGC	4188
2955	CAGCCTGG	GGCTAGCTACACGA	ACCTAAGC	3479	GCUUAGGU G CCAGGCUG	4189
2960	GCTTACAG	GGCTAGCTACACGA	CTGGCACC	3480	GGUGCCAG G CUGUAAGC	4190
2963	AATGCTTA	GGCTAGCTACACGA	AGCCTGGC	3481	GCCAGGCU G UAAGCAUU	4191
2967	TCAGAATG	GGCTAGCTACACGA	TTACAGCC	3482	GGCUGUA G CAUUCUGA	4192

Table 6

2969	GCTCAGAA GGCTAGCTACAACGA GCTTACAG	3483	CUGUAGC A UUCUGAGC	4193
2976	CAAGCCAG GGCTAGCTACAACGA TCAGAAATG	3484	CAUUCUGA G CUGGCUUG	4194
2980	ACAACAAG GGCTAGCTACAACGA CAGCTCAG	3485	CUGAGCUG G CUUGUUGU	4195
2984	AAAAACAA GGCTAGCTACAACGA AAGCCAGC	3486	GCUGGCUU G UUGUUUUU	4196
2987	CTTAAAAA GGCTAGCTACAACGA AACAGCC	3487	GGCUUGUU G UUUUUAAG	4197
2995	ATACAGGA GGCTAGCTACAACGA TTAAAAAC	3488	GUUUUAAA G UCCUGUAU	4198
3000	TACATATA GGCTAGCTACAACGA AGGACTTA	3489	UAAGUCCU G UAUUAUGA	4199
3002	CATACATA GGCTAGCTACAACGA ACAGGACT	3490	AGUCCUGU A UAUGUAUG	4200
3004	TACATACA GGCTAGCTACAACGA ATACAGGA	3491	UCCUGUAU A UGUUAUGA	4201
3006	ACTACATA GGCTAGCTACAACGA ATATACAG	3492	CUGUAUUA G UAUGUAGU	4202
3008	CTACTACA GGCTAGCTACAACGA ACATATAC	3493	GUUAUAUG A UGUUAUG	4203
3010	AACTACTA GGCTAGCTACAACGA ATACATAT	3494	AUAUGUAU G UAGUAGUU	4204
3013	CCAAACTA GGCTAGCTACAACGA TACATACA	3495	UGUAUGUA G UAGUUUGG	4205
3016	CACCCAAA GGCTAGCTACAACGA TACTACAT	3496	AUGUAUGA G UUUUGGUG	4206
3022	TATACACA GGCTAGCTACAACGA CCAAACTA	3497	UAGUUUGG G UGUUAUA	4207
3024	TATATACA GGCTAGCTACAACGA ACCCAAAC	3498	GUUUUGGU G UGUUAUA	4208
3026	TATATATA GGCTAGCTACAACGA ACACCCAA	3499	UUGGGUGU G UAUUAUA	4209
3028	ACTATATA GGCTAGCTACAACGA ACACACCC	3500	GGGUGUGU A UAUUAUGU	4210
3030	CTACTATA GGCTAGCTACAACGA ATACACAC	3501	GUGUGUAU A UAUAGUAG	4211
3032	TGCTACTA GGCTAGCTACAACGA ATATACAC	3502	GUGUAUAU A UAGUAGCA	4212
3035	AAATGCTA GGCTAGCTACAACGA TATATATA	3503	UAUAUAUA G UAGCAUUU	4213
3038	TTGAAATG GGCTAGCTACAACGA TACTATAT	3504	AUAUAGUA G CAUUUCA	4214
3040	TTTTGAAA GGCTAGCTACAACGA GCTACTAT	3505	AUAGUAGC A UUUCAA	4215
3048	TACGTCCA GGCTAGCTACAACGA TTGAAAT	3506	AUUUCAA A UGGACGUA	4216
3052	CCAGTACG GGCTAGCTACAACGA CCATTTTG	3507	CAAAUGG A CGUACUGG	4217
3054	AACCAGTA GGCTAGCTACAACGA GTCCATTT	3508	AAAUGGAC G UACUGGUU	4218
3056	TAAACCAG GGCTAGCTACAACGA ACCTCCAT	3509	AUGGACGU A CUGGUUUA	4219
3060	AGGTTAAA GGCTAGCTACAACGA CAGTACGT	3510	ACGUACUG G UUUAAACCU	4220
3065	ATAGGAGG GGCTAGCTACAACGA TAAACCCAG	3511	CUGGUUUA A CCUCCUAU	4221
3072	TCCAAGGA GGCTAGCTACAACGA AGGAGGTT	3512	AACCUCCU A UCCUUUGA	4222
3083	GCCAGCTG GGCTAGCTACAACGA TCTCCAAG	3513	CUUGGAGA G CAGCUGGC	4223

Table 6

3086	AGAGCCAG	GGCTAGCTACAACGA	TGCTCTCC	3514	GGAGAGCA	G CUGGCUCU	4224
3090	GTGGAGAG	GGCTAGCTACAACGA	CAGCTGCT	3515	AGCAGCUG	G CUCUCCAC	4225
3097	TAACAAGG	GGCTAGCTACAACGA	GGAGAGCC	3516	GGCUCUCC	A CCUUGUUA	4226
3102	ATGTGTAA	GGCTAGCTACAACGA	AAGGTGGA	3517	UCCACCUU	G UUACACAU	4227
3105	ATAATGTG	GGCTAGCTACAACGA	AACAAGGT	3518	ACCUUGUU	A CACAUUUA	4228
3107	ACATAATG	GGCTAGCTACAACGA	GTAACAAG	3519	CUUGUAC	A CAUUAUGU	4229
3109	TAACATTA	GGCTAGCTACAACGA	GTGTAACA	3520	UGUACAC	A UUAUGUUA	4230
3112	CTCTAACA	GGCTAGCTACAACGA	AATGTGTA	3521	UACACAUU	A UGUUAGAG	4231
3114	CTCTCTAA	GGCTAGCTACAACGA	ATAATGTG	3522	CACAUUUA	G UUAGAGAG	4232
3123	GCTCGCTA	GGCTAGCTACAACGA	CTCTCTAA	3523	UUAGAGAG	G UAGCGAGC	4233
3126	GCAGCTCG	GGCTAGCTACAACGA	TACCTCTC	3524	GAGAGGUA	G CGAGCUGC	4234
3130	CAGAGCAG	GGCTAGCTACAACGA	TCGCTACC	3525	GGUAGCGA	G CUGCUCUG	4235
3133	TAGCAGAG	GGCTAGCTACAACGA	AGCTCGCT	3526	AGCGAGCU	G CUCUGCUA	4236
3138	GGACATAG	GGCTAGCTACAACGA	AGAGCAGC	3527	GCUGCUCU	G CUUUGUCC	4237
3141	TAAGGACA	GGCTAGCTACAACGA	AGCAGAGC	3528	GCUCUGCU	A UGUCCUUA	4238
3143	CTTAAGGA	GGCTAGCTACAACGA	ATAGCAGA	3529	UCUGCUAU	G UCCUUAAG	4239
3151	AATATTGG	GGCTAGCTACAACGA	TAAAGGAC	3530	GUCUUAUA	G CCAAUUUA	4240
3155	AGTAAATA	GGCTAGCTACAACGA	TGGCTTAA	3531	UUAAGCCA	A UAUUUACU	4241
3157	TGAGTAA	GGCTAGCTACAACGA	ATTGGCTT	3532	AAGCCAAU	A UUUUACUA	4242
3161	CTGATGAG	GGCTAGCTACAACGA	AAATATTG	3533	CAAUUAUU	A CUCAUCAG	4243
3165	TGACCTGA	GGCTAGCTACAACGA	GAGTAAAT	3534	AUUUACUC	A UCAGGUCA	4244
3170	AATAATGA	GGCTAGCTACAACGA	CTGATGAG	3535	CUCAUCAG	G UCAUUAUU	4245
3173	AAAAATAA	GGCTAGCTACAACGA	GACCTGAT	3536	AUCAGGUC	A UUAUUUUU	4246
3176	GTAATAAA	GGCTAGCTACAACGA	AATGACCT	3537	AGGUCAUU	A UUUUUUAC	4247
3183	GGCCATTG	GGCTAGCTACAACGA	AAAAATAA	3538	UAUUUUUU	A CAUUGGCC	4248
3186	CATGGCCA	GGCTAGCTACAACGA	TGTAAAAA	3539	UUUUUACA	A UGGCCAUU	4249
3189	TTCCATGG	GGCTAGCTACAACGA	CATTGTAA	3540	UUACAAUG	G CCAUGGAA	4250
3192	TTATTCCA	GGCTAGCTACAACGA	GGCCATTG	3541	CAUUGGCC	A UGGAUAAA	4251
3197	ATGGTTTA	GGCTAGCTACAACGA	TCCATGGC	3542	GCCAUGGA	A UAAACCAU	4252
3201	AAAAATGG	GGCTAGCTACAACGA	TTATTCCA	3543	UGGAAUAA	A CCAUUUUU	4253
3204	TGTAAAAA	GGCTAGCTACAACGA	GGTTTATT	3544	AAUAAACC	A UUUUUACA	4254

Table 7

Table 7: Human PTP-1B Hairpin Ribozyme and Target Sequence

Nt. Position	Ribozyme sequence	Seq. ID Nos.	Substrate Sequence	Seq. ID Nos.
10	GCUCUA AGAA GCGU ACCAGAGAAAACA X GUACAUUACCUUGUA	4255	ACGC GGCC TAGAGC	4331
23	UGCGCC AGAA GCGC ACCAGAGAAAACA X GUACAUUACCUUGUA	4256	CGGC AGAC GCGCA	4332
56	CAGGGC AGAA GCUG ACCAGAGAAAACA X GUACAUUACCUUGUA	4257	CAGC AGCC GCGCTG	4333
59	GGCCAG AGAA GCUG ACCAGAGAAAACA X GUACAUUACCUUGUA	4258	CAGC CGCC CTGGCC	4334
98	UUGUCG AGAA GCUC ACCAGAGAAAACA X GUACAUUACCUUGUA	4259	GAGC AGAT CGACAA	4335
220	CAAAAG AGAA GACG ACCAGAGAAAACA X GUACAUUACCUUGUA	4260	CGTC AGTC CCTTGG	4336
239	AGUUUA AGAA GACU ACCAGAGAAAACA X GUACAUUACCUUGUA	4261	AGTC GGAT TAAACT	4337
612	UCCAAA AGAA GGCC ACCAGAGAAAACA X GUACAUUACCUUGUA	4262	GGCC TGAC TTTGGA	4338
636	GAAUGA AGAA GGUG ACCAGAGAAAACA X GUACAUUACCUUGUA	4263	CACC AGCC TCATTC	4339
685	GCUCGC AGAA GAGU ACCAGAGAAAACA X GUACAUUACCUUGUA	4264	ACTC AGCC CGGAGC	4340
702	CACCAC AGAA GGCC ACCAGAGAAAACA X GUACAUUACCUUGUA	4265	GGCC CGTT GTGGTG	4341
748	CAGCCA AGAA GAAG ACCAGAGAAAACA X GUACAUUACCUUGUA	4266	CTTC TGTC TGGCTG	4342
763	GCAAGA AGAA GGUU ACCAGAGAAAACA X GUACAUUACCUUGUA	4267	TACC TGCC TCTTGC	4343
773	UUGUCC AGAA GCAA ACCAGAGAAAACA X GUACAUUACCUUGUA	4268	TTGC TGAT GGACAA	4344
801	GAUAUC AGAA GAAG ACCAGAGAAAACA X GUACAUUACCUUGUA	4269	CTTC CGTT GATATC	4345
842	AGCCCC AGAA GAAA ACCAGAGAAAACA X GUACAUUACCUUGUA	4270	TTTC GGAT GGGGCT	4346
851	GUUCUG AGAA GCCC ACCAGAGAAAACA X GUACAUUACCUUGUA	4271	GGGC TGAT CCAGAC	4347
861	CUUGUC AGAA GUCU ACCAGAGAAAACA X GUACAUUACCUUGUA	4272	AGAC AGCC GACCAG	4348
864	CAGCUG AGAA GCUG ACCAGAGAAAACA X GUACAUUACCUUGUA	4273	CAGC CGAC CAGCTG	4349
869	AAGGCG AGAA GGUC ACCAGAGAAAACA X GUACAUUACCUUGUA	4274	GACC AGCT GCGCTT	4350
1102	UGAUGG AGAA GUCU ACCAGAGAAAACA X GUACAUUACCUUGUA	4275	AGAC TGCC CCATCA	4351
1224	UGGGGA AGAA GCCU ACCAGAGAAAACA X GUACAUUACCUUGUA	4276	AGGC TGCC TCCCCA	4352
1253	UUCUCG AGAA GUGA ACCAGAGAAAACA X GUACAUUACCUUGUA	4277	TCAC TGCC CGAGAA	4353
1323	CGUGAG AGAA GUAG ACCAGAGAAAACA X GUACAUUACCUUGUA	4278	CTAC GGTC CTCACG	4354
1332	AGCGCC AGAA GUGA ACCAGAGAAAACA X GUACAUUACCUUGUA	4279	TCAC GGCC GGCCT	4355
1361	CUGUUG AGAA GGAA ACCAGAGAAAACA X GUACAUUACCUUGUA	4280	TTCC TGTG CAACAG	4356
1383	AGGAGG AGAA GGCU ACCAGAGAAAACA X GUACAUUACCUUGUA	4281	AGCC TGAC CCTCCT	4357
1412	GAGGCG AGAA GUGG ACCAGAGAAAACA X GUACAUUACCUUGUA	4282	CCAC TGTC CGCCTC	4358

Table 7

1416	GGCAGA AGAA GACA ACCAGAGAAACA X GUACAUUACCUUGUA	4283	TGTC CGCC TCTGCC	4359
1422	UCUGCG AGAA GAGG ACCAGAGAAACA X GUACAUUACCUUGUA	4284	CCTC TGCC CGCAGA	4360
1441	CUGCUA AGAA GCGC ACCAGAGAAACA X GUACAUUACCUUGUA	4285	CGCC CGAC TAGCAG	4361
1460	CUUACC AGAA GCGG ACCAGAGAAACA X GUACAUUACCUUGUA	4286	CCGC GGTA GGTAAG	4362
1473	CGGUCC AGAA GCCC ACCAGAGAAACA X GUACAUUACCUUGUA	4287	GGGC CGCC GGACCG	4363
1477	UACGCG AGAA GCGC ACCAGAGAAACA X GUACAUUACCUUGUA	4288	CGCC GGAC CGCGTA	4364
1501	ACGUCC AGAA GGGG ACCAGAGAAACA X GUACAUUACCUUGUA	4289	CCCC GGAC GGACGT	4365
1505	ACCAAC AGAA GUCC ACCAGAGAAACA X GUACAUUACCUUGUA	4290	GGAC GGAC GTTGGT	4366
1539	ACACAC AGAA GGGG ACCAGAGAAACA X GUACAUUACCUUGUA	4291	CCCC GGAT GTGTGT	4367
1670	CCCCAA AGAA GGUG ACCAGAGAAACA X GUACAUUACCUUGUA	4292	CACC CGTC TTGGGG	4368
1735	GGGGGG AGAA GUCU ACCAGAGAAACA X GUACAUUACCUUGUA	4293	AGAC AGCC CCCCCC	4369
1864	AUGCAA AGAA GACA ACCAGAGAAACA X GUACAUUACCUUGUA	4294	TGTC AGCC TTGCAT	4370
1946	UGACGC AGAA GGCC ACCAGAGAAACA X GUACAUUACCUUGUA	4295	GGCC TGCT GCGTCA	4371
1955	GUACUG AGAA GACG ACCAGAGAAACA X GUACAUUACCUUGUA	4296	CGTC AGAC CAGTAC	4372
1960	UCCACG AGAA GGUC ACCAGAGAAACA X GUACAUUACCUUGUA	4297	GACC AGTA CTGGGA	4373
1978	GCUIAC AGAA GUCC ACCAGAGAAACA X GUACAUUACCUUGUA	4298	GGAC GGTG GTNAGC	4374
1988	AUAUAC AGAA GCUU ACCAGAGAAACA X GUACAUUACCUUGUA	4299	AAGC AGTT GTTATT	4375
2098	AGAAUA AGAA GGAC ACCAGAGAAACA X GUACAUUACCUUGUA	4300	GTCC CGCT TATTCT	4376
2107	ACAGGG AGAA GAAU ACCAGAGAAACA X GUACAUUACCUUGUA	4301	ATTG TGCT CCTGTG	4377
2144	CGGGGG AGAA GUGA ACCAGAGAAACA X GUACAUUACCUUGUA	4302	TCAC TGCT CCCCCG	4378
2190	UUUUUC AGAA GGAC ACCAGAGAAACA X GUACAUUACCUUGUA	4303	GTCC TGAT GAAAAA	4379
2230	UUAGUA AGAA GAGA ACCAGAGAAACA X GUACAUUACCUUGUA	4304	TCTC TGCT TACTAA	4380
2264	UGCACA AGAA GGUU ACCAGAGAAACA X GUACAUUACCUUGUA	4305	AACC TGCC TGTGCA	4381
2279	GUUAUG AGAA GGUC ACCAGAGAAACA X GUACAUUACCUUGUA	4306	GACC TGAT CATTAC	4382
2309	UUCAGC AGAA GCGU ACCAGAGAAACA X GUACAUUACCUUGUA	4307	AGCC TGTT GCTGAA	4383
2345	CUGGAA AGAA GCAC ACCAGAGAAACA X GUACAUUACCUUGUA	4308	GTGC AGTT TTCCAG	4384
2412	AGGACA AGAA GGGC ACCAGAGAAACA X GUACAUUACCUUGUA	4309	GCCC AGCC TGTCTT	4385
2416	UACCAG AGAA GCGU ACCAGAGAAACA X GUACAUUACCUUGUA	4310	AGCC TGTC CTGGTA	4386
2524	GUGGGG AGAA GGGG ACCAGAGAAACA X GUACAUUACCUUGUA	4311	CCCC CGCC CCCCAC	4387
2544	GGCGGA AGAA GAUC ACCAGAGAAACA X GUACAUUACCUUGUA	4312	GATC AGCC TCCGCC	4388
2580	AUCACG AGAA GCUC ACCAGAGAAACA X GUACAUUACCUUGUA	4313	GAGC AGAC CGTGAT	4389

Table 7

2606	GUUCC AGAA GGUG ACCAGAGAAACA X GUACAUUACCUUGUA	4314	CACC TGCT GGAAC	4390
2631	CACCCA AGAA GUUU ACCAGAGAAACA X GUACAUUACCUUGUA	4315	AAAC AGCC TGGTG	4391
2643	CUAAAG AGAA GUCA ACCAGAGAAACA X GUACAUUACCUUGUA	4316	TGAC GGTC CTTTAG	4392
2655	GCGCA AGAA GCCU ACCAGAGAAACA X GUACAUUACCUUGUA	4317	AGGC AGCC TGCOC	4393
2659	GACGC AGAA GGCU ACCAGAGAAACA X GUACAUUACCUUGUA	4318	AGCC TGCC GCCGTC	4394
2662	AGAGAC AGAA GCAG ACCAGAGAAACA X GUACAUUACCUUGUA	4319	CTGC CGCC GTCTCT	4395
2665	GACAGA AGAA GCGG ACCAGAGAAACA X GUACAUUACCUUGUA	4320	CCGC CGTC TCTGTC	4396
2671	AACCGG AGAA GAGA ACCAGAGAAACA X GUACAUUACCUUGUA	4321	TCTC TGTC CCGGTT	4397
2677	AAGGUG AGAA GGA ACCAGAGAAACA X GUACAUUACCUUGUA	4322	TCCC GGT CACCTT	4398
2704	GGUGG AGAA GACG ACCAGAGAAACA X GUACAUUACCUUGUA	4323	CGTC TGCC CCACCC	4399
2731	AGCAC AGAA GGCC ACCAGAGAAACA X GUACAUUACCUUGUA	4324	GGCC TGAT GGTGCT	4400
2810	AGCUAC AGAA GCCG ACCAGAGAAACA X GUACAUUACCUUGUA	4325	CGGC AGCT GTAGCT	4401
2813	GGGAGC AGAA GCUG ACCAGAGAAACA X GUACAUUACCUUGUA	4326	CAGC TGT GCTCCC	4402
2876	UCUCUG AGAA GGCU ACCAGAGAAACA X GUACAUUACCUUGUA	4327	AGCC AGTA CAGAGA	4403
2999	UACAU AGAA GGAC ACCAGAGAAACA X GUACAUUACCUUGUA	4328	GTCC TGT TATGTA	4404
3085	AGAGCC AGAA GCUC ACCAGAGAAACA X GUACAUUACCUUGUA	4329	GAGC AGCT GGCTCT	4405
3132	UAGCAG AGAA GCUC ACCAGAGAAACA X GUACAUUACCUUGUA	4330	GAGC TGCT CTGCTA	4406

Table 8

Table 8: Anti Human PTP-1B HH, NCH, and G Cleaver Ribozymes

Alias	Ribozyme Sequence	Seq. ID Nos	Substrate Seq.	Seq. ID Nos.
HH				
PTP1B-599	UGUGUA CUGAUGAGGCCGUUAGGCCGAA AGUGGAA	4407	UUCCACU A UACCACA	4425
PTP1B-879	GGUAGGA CUGAUGAGGCCGUUAGGCCGAA AAGCGCA	4408	UGCGCUU C UCCUACC	4426
PTP1B-1393	UGGAGUG CUGAUGAGGCCGUUAGGCCGAA AGGAGGG	4409	CCCUCCU C CACUCCA	4427
PTP1B-1398	GGAGGUG CUGAUGAGGCCGUUAGGCCGAA AGUGGAG	4410	CUCCACU C CACCUCC	4428
PTP1B-1404	GUGGGUG CUGAUGAGGCCGUUAGGCCGAA AGUGGA	4411	UCCACCU C CACCCAC	4429
PTP1B-2118	UAGCAGA CUGAUGAGGCCGUUAGGCCGAA AACAGGG	4412	CCCUUUU A UCUGCUA	4430
PTP1B-2181	GACACAA CUGAUGAGGCCGUUAGGCCGAA AAGACCU	4413	AGGUCUU C UUGUGUC	4431
PTP1B-2183	AGGACAC CUGAUGAGGCCGUUAGGCCGAA AGAAGAC	4414	GUCUUUU U GUGUCCU	4432
PTP1B-2238	GGCACAU CUGAUGAGGCCGUUAGGCCGAA AGUAAGC	4415	GCUUACU A AUGUGCC	4433
PTP1B-2252	GGACUUG CUGAUGAGGCCGUUAGGCCGAA ACAUGGG	4416	CCCAUGU C CAAGUCC	4434
NCH				
PTP1B-1395	GGUGGAG CUGAUGAGGCCGUUAGGCCGAA IGAGGAG	4417	CUCCUCC A CUCCACC	4435
PTP1B-1408	GACAGUG CUGAUGAGGCCGUUAGGCCGAA IUGGAGG	4418	CCUCCAC C CACUGUC	4436
PTP1B-1555	GGAUGAG CUGAUGAGGCCGUUAGGCCGAA IGUGAGA	4419	UCUCACC C CUCAUCC	4437
PTP1B-1578	AGUGGAA CUGAUGAGGCCGUUAGGCCGAA IGGCAAA	4420	UUUGCCC C UUCCACU	4438
PTP1B-2113	GAUAACA CUGAUGAGGCCGUUAGGCCGAA IGAGCAG	4421	CUGCUCC C UGUUAUC	4439
G-Cleaver				
PTP1B-1267	GUCCU UGAUGGCAUGCACUAUGCCGG GUCCUUCUG	4422	CGAGAAGGAC G AGGAC	4440
PTP1B-2184	GGACA UGAUGGCAUGCACUAUGCCGG AAGAAGACCU	4423	AGGUUUUUU G UGUCC	4441
PTP1B-2241	GGGCA UGAUGGCAUGCACUAUGCCGG AUUAGUAAGC	4424	GCUUACUAAU G UGCCC	4442

Table 9

Table 9: Human methionine aminopeptidase type 2 (Met AP-2) Hammerhead Ribozyme and Target Sequence

Nt. position	Ribozyme Sequence	Seq ID nos.	Substrate Sequence	Seq ID nos.
9	CCGAGAGA CUGAUGAG X CGAA ACGAGGGA	1	TCCCTCGT C TCTCTCGG	413
11	GCCCGAGA CUGAUGAG X CGAA AGACGAGG	2	CCTCGTCT C TCTCGGGC	414
13	UUGCCCGA CUGAUGAG X CGAA AGAGACGA	3	TCGTCTCT C TCGGGCAA	415
15	UGUUGCCC CUGAUGAG X CGAA AGAGAGAC	4	GTCTCTCT C GGGCAACA	416
43	GAGGCCGC CUGAUGAG X CGAA ACCUCCUC	5	GAGGAGGT A GCGGCCTC	417
51	GGCUCCCG CUGAUGAG X CGAA AGGCCGCU	6	AGCGGCCT C CGGGAGCC	418
80	GUCGUCUG CUGAUGAG X CGAA AUCCAGGU	7	ACCTGGAT C CAGACGAC	419
108	CAGCCGUA CUGAUGAG X CGAA AGGCAGCU	8	AGCTGCCT C TACGGCTG	420
110	CUCAGCCG CUGAUGAG X CGAA AGAGGCAG	9	CTGCCTCT A CGGCTGAG	421
167	UGCUGCAG CUGAUGAG X CGAA AGGCCCUU	10	AAGGGCCT T CTCGAGCA	422
168	CUGCUGCA CUGAUGAG X CGAA AAGGCCCU	11	AGGGCCTT C TGCAGCAG	423
194	UGAUUCUU CUGAUGAG X CGAA AUCAGGUU	12	AACCTGAT A AAGAATCA	424
201	AGGCUCCU CUGAUGAG X CGAA AUUCUUUA	13	TAAAGAAAT C AGGAGCCT	425
210	CAUCCACU CUGAUGAG X CGAA AGGCUCCU	14	AGGAGCCT C AGTGGATG	426
223	UGUCUUGC CUGAUGAG X CGAA ACUUCAUC	15	GATGAAGT A GCAAGACA	427
234	AUCUUUCC CUGAUGAG X CGAA ACUGUCUU	16	AAGACAGT T GGAAAGAT	428
243	CCAAUGCU CUGAUGAG X CGAA AUCUUUCC	17	GGAAAGAT C AGCATTGG	429
249	UAUCUUCC CUGAUGAG X CGAA AUGCUGAU	18	ATCAGCAT T GGAAGATA	430
257	UCUUUCUU CUGAUGAG X CGAA AUCUCCA	19	TGGAAGAT A AAGAAAGA	431
355	UCUGUUUG CUGAUGAG X CGAA ACUUUUGG	20	CCAAAAGT T CAAACAGA	432
356	GUCUGUUU CUGAUGAG X CGAA AACUUUUG	21	CAAAAGTT C AAACAGAC	433
368	AACUGAGG CUGAUGAG X CGAA AGGGUCUG	22	CAGACCCT C CCTCAGTT	434
372	UUGGAACU CUGAUGAG X CGAA AGGGAGGG	23	CCCTCCCT C AGTTCCAA	435
376	CAUAUUGG CUGAUGAG X CGAA ACUGAGGG	24	CCCTCAGT T CCAATATG	436
377	ACAUAUUG CUGAUGAG X CGAA AACUGAGG	25	CCTCAGTT C CAATATGT	437
382	AGGUCACA CUGAUGAG X CGAA AUUGGAAC	26	GTTCCAAT A TGTGACCT	438
393	CAUUAAGG CUGAUGAG X CGAA ACAGGUCA	27	TGACCTGT A TCCTAATG	439
395	ACCAUUAG CUGAUGAG X CGAA AUACAGGU	28	ACCTGTAT C CTAATGGT	440
398	UACACCAU CUGAUGAG X CGAA AGGAUACA	29	TGTATCCT A ATGGTGTA	441
406	UUGGGAAA CUGAUGAG X CGAA ACACCAUU	30	AATGGTGT A TTTCCTAA	442
408	CUUUGGGA CUGAUGAG X CGAA AUACACCA	31	TGGTGTAT T TCCCAAAG	443
409	CCUUGGGG CUGAUGAG X CGAA AAUACACC	32	GGTGTATT T CCCAAAGG	444
410	UCCUUUGG CUGAUGAG X CGAA AAUACAC	33	GTGTATTT C CCAAAGGA	445
432	UGGGUGGG CUGAUGAG X CGAA AUUCGCAU	34	ATGCGAAT A CCCACCCA	446
464	AGUUCUCC CUGAUGAG X CGAA AGCAGCUG	35	CAGCTGCT T GGAGAACT	447
473	UUCACUUG CUGAUGAG X CGAA AGUUCUCC	36	GGAGAACT A CAAGTGAA	448
495	CCUGAUCU CUGAUGAG X CGAA AUGCUUUC	37	GAAAGCAT T AGATCAGG	449
496	GCCUGAUC CUGAUGAG X CGAA AAUGCUUU	38	AAAGCATT A GATCAGGC	450
500	ACUUGCCU CUGAUGAG X CGAA AUCUAAUG	39	CATTAGAT C AGGCAAGT	451
517	UCAUUGCA CUGAUGAG X CGAA AUCUCUUC	40	GAAGAGAT T TGAATGTA	452
518	AUCAUUC CUGAUGAG X CGAA AAUCUCUU	41	AAGAGATT T GGAATGAT	453
527	UUCUCGAA CUGAUGAG X CGAA AUCAUUC	42	GGAATGAT T TTCGAGAA	454

Table 9

528	CUUCUCGA CUGAUGAG X CGAA AAUCAUUC	43	GAATGATT T TCGAGAAG	455
529	GCUUCUCG CUGAUGAG X CGAA AAAUCAU	44	AATGATTT T CGAGAAGC	456
530	AGCUUCUC CUGAUGAG X CGAA AAAUCAU	45	ATGATTTT C GAGAAGCT	457
551	AACUUGUC CUGAUGAG X CGAA AUGUGCUU	46	AAGCACAT C GACAAGTT	458
559	UAUUUUCU CUGAUGAG X CGAA ACUUGUCG	47	CGACAAGT T AGAAAATA	459
560	GUUUUUC CUGAUGAG X CGAA AACUUGUC	48	GACAAGTT A GAAAATAC	460
567	UCAUUACG CUGAUGAG X CGAA AUUUUCUA	49	TAGAAAAT A CGTAATGA	461
571	CAGCUCAU CUGAUGAG X CGAA ACGUAUUU	50	AAATACGT A ATGAGCTG	462
583	CCAGGCUU CUGAUGAG X CGAA AUCCAGCU	51	AGCTGGAT C AAGCCTGG	463
604	CAGAUUUC CUGAUGAG X CGAA AUCAUUGU	52	ACAATGAT A GAAATCTG	464
610	UUUUCACA CUGAUGAG X CGAA AUUUCUAU	53	ATAGAAAT C TGTGAAAA	465
621	AGUCUUC CUGAUGAG X CGAA ACUUUUCA	54	TGAAAAGT T GGAAGACT	466
632	CUUGCGUG CUGAUGAG X CGAA ACAGUCUU	55	AAGACTGT T CACGCAAG	467
633	ACUUGCGU CUGAUGAG X CGAA AACAGUCU	56	AGACTGTT C ACGCAAGT	468
642	CUUUUAUU CUGAUGAG X CGAA ACUUGCGU	57	ACGCAAGT T AATAAAG	469
643	UCUUUUAU CUGAUGAG X CGAA AACUUGCG	58	CGCAAGTT A ATAAAAGA	470
646	UUCUCUUU CUGAUGAG X CGAA AUUAACUU	59	AAGTTAAT A AAAGAGAA	471
660	CUGCAUUU CUGAUGAG X CGAA AUCCAUUC	60	GAATGGAT T AAATGCAG	472
661	CCUGCAUU CUGAUGAG X CGAA AAUCCAUU	61	AATGGATT A AATGCAGG	473
678	CAGUAGGA CUGAUGAG X CGAA AUGCCAGG	62	CCTGGCAT T TCCTACTG	474
679	CCAGUAGG CUGAUGAG X CGAA AAUGCCAG	63	CTGGCATT T CCTACTGG	475
680	UCCAGUAG CUGAUGAG X CGAA AAAUGCCA	64	TGGCATT T C TACTGGA	476
683	ACAUCCAG CUGAUGAG X CGAA AGGAAAUG	65	CATTTCCT A CTGGATGT	477
692	AUUGAGAG CUGAUGAG X CGAA ACAUCCAG	66	CTGGATGT T CTCTCAAT	478
693	UAUUGAGA CUGAUGAG X CGAA AACAUCCA	67	TGGATGTT C TCTCAATA	479
695	AUUUAUGA CUGAUGAG X CGAA AGAACAUC	68	GATGTCTCT C TCAATAAT	480
697	CAAUAUUU CUGAUGAG X CGAA AGAGAACA	69	TGTTCTCT C AATAATTG	481
701	AGCACA AU CUGAUGAG X CGAA AUUGAGAG	70	CTCTCAAT A ATTGTGCT	482
704	GGCAGCAC CUGAUGAG X CGAA AUUAUUGA	71	TCAATAAT T GTGCTGCC	483
716	GGGAGUAU CUGAUGAG X CGAA AUGGGCAG	72	CTGCCCAT T ATACTCCC	484
717	UGGGAGUA CUGAUGAG X CGAA AAUGGGCA	73	TGCCCCATT A TACTCCCA	485
719	AUUGGGAG CUGAUGAG X CGAA AUAAUGGG	74	CCCATTAT A CTCCCAAT	486
722	GGCAUUGG CUGAUGAG X CGAA AGUAUAAU	75	ATTATACT C CCAATGCC	487
745	UACUGUAA CUGAUGAG X CGAA ACUGUUGU	76	ACAACAGT A TTACAGTA	488
747	CAUACUGU CUGAUGAG X CGAA AUACUGUU	77	AACAGTAT T ACAGTATG	489
748	UCAUACUG CUGAUGAG X CGAA AAUACUGU	78	ACAGTATT A CAGTATGA	490
753	UGUCAUCA CUGAUGAG X CGAA ACUGUAAU	79	ATTACAGT A TGATGACA	491
763	AUUUUAACA CUGAUGAG X CGAA AUGUCAUC	80	GATGACAT C TGTAATAAT	492
767	GUCUAUUU CUGAUGAG X CGAA ACAGAUGU	81	ACATCTGT A AAATAGAC	493
772	CCAAAGUC CUGAUGAG X CGAA AUUUUACA	82	TGTAAAAT A GACTTTGG	494
777	GUGUUGCA CUGAUGAG X CGAA AGUCUAUU	83	AATAGACT T TGAACAC	495
778	UGUGUUC CUGAUGAG X CGAA AAGUCUAU	84	ATAGACTT T GGAACACA	496
788	ACCACUUA CUGAUGAG X CGAA AUGUGUUC	85	GAACACAT A TAAGTGGT	497
790	CUACCACU CUGAUGAG X CGAA AUAUGUGU	86	ACACATAT A AGTGGTAG	498
797	AAUAAUCC CUGAUGAG X CGAA ACCACUUA	87	TAAGTGGT A GGATTATT	499
802	CAGUCAAU CUGAUGAG X CGAA AUCCUACC	88	GGTAGGAT T ATTGACTG	500
803	ACAGUCAA CUGAUGAG X CGAA AAUCCUAC	89	GTAGGATT A TTGACTGT	501

Table 9

805	GCACAGUC CUGAUGAG X CGAA AUAUCCU	90	AGGATTAT T GACTGTGC	502
815	GACAGUAA CUGAUGAG X CGAA AGCACAGU	91	ACTGTGCT T TACTGTGC	503
816	UGACAGUA CUGAUGAG X CGAA AAGCACAG	92	CTGTGCTT T TACTGTCA	504
817	GUGACAGU CUGAUGAG X CGAA AAAGCACA	93	TGTGCTTT T ACTGTCACT	505
818	AGUGACAG CUGAUGAG X CGAA AAAAGCAC	94	GTGCTTTT A CTGTCACT	506
823	UUAAAAGU CUGAUGAG X CGAA ACAGUAAA	95	TTTACTGT C ACTTTTAA	507
827	GGGAUUA CUGAUGAG X CGAA AGUGACAG	96	CTGTCACT T TTAATCCC	508
828	UGGGAUUA CUGAUGAG X CGAA AAGUGACA	97	TGTCACTT T TAATCCCA	509
829	UUGGGAUU CUGAUGAG X CGAA AAAGUGAC	98	GTCACCTT T AATCCCAA	510
830	UUUGGGAU CUGAUGAG X CGAA AAAAGUGA	99	TCACTTTT A ATCCCAAA	511
833	AUAUUUGG CUGAUGAG X CGAA AUUAAAAG	100	CTTTTAAT C CCAAATAT	512
840	ACGUAUCA CUGAUGAG X CGAA AUUUGGGA	101	TCCCAAAAT A TGATACGT	513
845	UAAUAACG CUGAUGAG X CGAA AUCAUAUU	102	AATATGAT A CGTTATTA	514
849	CUUUAAU CUGAUGAG X CGAA ACGUAUCA	103	TGATACGT T ATTAAGAAG	515
850	GCUUUUAA CUGAUGAG X CGAA AACGUUUC	104	GATACGTT A TTAAGAAGC	516
852	CAGCUUUU CUGAUGAG X CGAA AUAACGUA	105	TACGTAT T AAAAGCTG	517
853	ACAGCUUU CUGAUGAG X CGAA AAUAACGU	106	ACGTTATT A AAAGCTGT	518
862	GCAUCUUU CUGAUGAG X CGAA ACAGCUUU	107	AAAGCTGT A AAAGATGC	519
872	AGUGUUAG CUGAUGAG X CGAA AGCAUCUU	108	AAGATGCT A CTAACACT	520
875	UCCAGUGU CUGAUGAG X CGAA AGUAGCAU	109	ATGCTACT A AACTGGA	521
886	GCACACUU CUGAUGAG X CGAA AUUCCAGU	110	ACTGGAAT A AAGTGTGC	522
901	CGAACAU CUGAUGAG X CGAA AUUCCAGC	111	GCTGGAAT T GATGTTCC	523
907	CACAGACG CUGAUGAG X CGAA ACAUCAAU	112	ATTGATGT T CGTCTGTG	524
908	ACACAGAC CUGAUGAG X CGAA AACAUCAA	113	TTGATGTT C GTCTGTGT	525
911	AUCACACA CUGAUGAG X CGAA ACGAACAU	114	ATGTTCTG C TGTGTGAT	526
922	GCCUCACC CUGAUGAG X CGAA ACAUCACA	115	TGTGATGT T GGTGAGGC	527
934	ACUUCUUG CUGAUGAG X CGAA AUGGCCUC	116	GAGGCCAT C CAAGAAGT	528
943	GACUCCA CUGAUGAG X CGAA ACUUCUUG	117	CAAGAAGT T ATGGAGTC	529
944	GGACUCCA CUGAUGAG X CGAA AACUUCUU	118	AAGAAGTT A TGGAGTCC	530
951	CUUCAUAG CUGAUGAG X CGAA ACUCCAUA	119	TATGGAGT C CTATGAAG	531
954	CAACUUCA CUGAUGAG X CGAA AGGACUCC	120	GGAGTCCT A TGAAGTTG	532
961	UCUAUUUC CUGAUGAG X CGAA ACUUCAUA	121	TATGAAGT T GAAATAGA	533
967	UUCCCAUC CUGAUGAG X CGAA AUUUCAAC	122	GTTGAAAT A GATGGGAA	534
981	UCACUUGA CUGAUGAG X CGAA AUGUCUUC	123	GAAGACAT A TCAAGTGA	535
983	UUUCACUU CUGAUGAG X CGAA AUAUGUCU	124	AGACATAT C AAGTGAAA	536
997	AGAUUACG CUGAUGAG X CGAA AUUGGUUU	125	AAACCAAT C CGTAATCT	537
1001	AUUUAGAU CUGAUGAG X CGAA ACGAUUG	126	CAATCCGT A ATCTAAAT	538
1004	UCCAUUUA CUGAUGAG X CGAA AUUACGGA	127	TCCGTAAT C TAAATGGA	539
1006	UGUCCAUA CUGAUGAG X CGAA AGAUUACG	128	CGTAATCT A AATGGACA	540
1016	CCCAAUUG CUGAUGAG X CGAA AUGUCCA	129	ATGGACAT T CAATTGGG	541
1017	GCCCAAUU CUGAUGAG X CGAA AAUGUCCA	130	TGGACATT C AATTGGGC	542
1021	UAUUGCCC CUGAUGAG X CGAA AUUGAAUG	131	CATTCAAT T GGGCAATA	543
1029	GUUUUUA CUGAUGAG X CGAA AUUGCCCA	132	TGGGCAAT A TAGAATAC	544
1031	AUGUAUUC CUGAUGAG X CGAA AUAUUGCC	133	GGCAATAT A GAATACAT	545
1036	CCAGCAUG CUGAUGAG X CGAA AUUCUAUA	134	TATAGAAT A CATGCTGG	546
1060	CCUUUCAC CUGAUGAG X CGAA AUCGGCAC	135	GTGCCGAT T GTGAAAGG	547
1102	AUUGCAUA CUGAUGAG X CGAA ACUUCUCC	136	GGAGAAGT A TATGCAAT	548

Table 9

1104	CAAUUGCA CUGAUGAG X CGAA AUACUUCU	137	AGAAGTAT A TGCAATTG	549
1111	AAGGUUUC CUGAUGAG X CGAA AUUGCAUA	138	TATGCAAT T GAAACCTT	550
1119	UACUACCA CUGAUGAG X CGAA AGGUUUCA	139	TGAAACCT T TGGTAGTA	551
1120	GUACUACC CUGAUGAG X CGAA AAGGUUUC	140	GAAACCTT T GGTAGTAC	552
1124	UCCUGUAC CUGAUGAG X CGAA ACCAAAGG	141	CCTTGGT A GTACAGGA	553
1127	UUUUCUG CUGAUGAG X CGAA ACUACCAA	142	TTGGTAGT A CAGGAAAA	554
1141	UCAUGAAC CUGAUGAG X CGAA ACACCUU	143	AAAGGTGT T GTTCATGA	555
1144	UCAUCAUG CUGAUGAG X CGAA ACAACACC	144	GGTGTGT T CATGATGA	556
1145	AUCAUCAU CUGAUGAG X CGAA AACAACAC	145	GTGTGTGT C ATGATGAT	557
1154	ACAUUCCA CUGAUGAG X CGAA AUCAUCAU	146	ATGATGAT A TGGAATGT	558
1163	GUAAUGUG CUGAUGAG X CGAA ACAUUCCA	147	TGGAATGT T CACATTAC	559
1164	UGUAAUGU CUGAUGAG X CGAA AACAUUCC	148	GGAATGTT C ACATTACA	560
1169	UUUCAUGU CUGAUGAG X CGAA AUGUGAAC	149	GTTACAT T ACATGAAA	561
1170	UUUCAUG CUGAUGAG X CGAA AAUGUGAA	150	TTCACATT A CATGAAAA	562
1181	AACAUCAA CUGAUGAG X CGAA AUUUUUA	151	TGAAAAAT T TTGATGTT	563
1182	CAACAUCA CUGAUGAG X CGAA AAUUUUUC	152	GAAAAATT T TGATGTTG	564
1183	CCAACAUC CUGAUGAG X CGAA AAAUUUUU	153	AAAAATTT T GATGTTGG	565
1189	ACAUGUCC CUGAUGAG X CGAA ACAUCAAA	154	TTTGATGT T GGACATGT	566
1204	GGAAGCCU CUGAUGAG X CGAA AUUGGCAC	155	GTGCCAAT A AGGCTTCC	567
1210	GUUCUUGG CUGAUGAG X CGAA AGCCUUAU	156	ATAAGGCT T CCAAGAAC	568
1211	UGUUCUUG CUGAUGAG X CGAA AAGCCUUA	157	TAAGGCTT C CAAGAACA	569
1227	CAUUAAC CUGAUGAG X CGAA AGUGUUUU	158	AAAACACT T GTTAAATG	570
1230	UGACAUUU CUGAUGAG X CGAA ACAAGUGU	159	ACACTTGT T AAATGTCA	571
1231	AUGACAUU CUGAUGAG X CGAA AACAAGUG	160	CACTTGTT A AATGTCAT	572
1237	UCAUUGAU CUGAUGAG X CGAA ACAUUUAA	161	TTAAATGT C ATCAATGA	573
1240	UUUCAUU CUGAUGAG X CGAA AUGACAUU	162	AATGTCAT C AATGAAAA	574
1251	GGGUUCCA CUGAUGAG X CGAA AGUUUUA	163	TGAAACT T TGAACCC	575
1252	AGGUUCC CUGAUGAG X CGAA AAGUUUUC	164	GAAAACTT T GGAACCTT	576
1261	CAGAAGGC CUGAUGAG X CGAA AGGUUCC	165	GGAACCTT T GCCTTCTG	577
1266	UGCGGCAG CUGAUGAG X CGAA AGGCAAGG	166	CCTTGCCT T CTGCCGCA	578
1267	CUGCGGCA CUGAUGAG X CGAA AAGGCAAG	167	CTTGCCTT C TGCCGAG	579
1286	UCCCAAGC CUGAUGAG X CGAA AUCCAGCC	168	GGCTGGAT C GCTTGGGA	580
1290	UUUCUCCC CUGAUGAG X CGAA AGCGAUCC	169	GGATCGCT T GGGAGAAA	581
1301	CAAGUAUU CUGAUGAG X CGAA ACUUUCUC	170	GAGAAAGT A AATACTTG	582
1305	CCAUCAAG CUGAUGAG X CGAA AUUUACUU	171	AAGTAAAT A CTTGATGG	583
1308	GAGCAUC CUGAUGAG X CGAA AGUAUUUA	172	TAAATACT T GATGGCTC	584
1316	AUUCUUA CUGAUGAG X CGAA AGCCAUA	173	TGATGGCT C TGAAGAAT	585
1325	GUCACACA CUGAUGAG X CGAA AUUCUUA	174	TGAAGAAT C TGTGTGAC	586
1335	CAAUGCCC CUGAUGAG X CGAA AGUCACAC	175	GTGTGACT T GGGCATTG	587
1342	GGAUCUAC CUGAUGAG X CGAA AUGCCCAA	176	TTGGGCAT T GTAGATCC	588
1345	UAUGGAUC CUGAUGAG X CGAA ACAUUGCC	177	GGCATTGT A GATCCATA	589
1349	UGGAUAUG CUGAUGAG X CGAA AUCUACAA	178	TTGTAGAT C CATATCCA	590
1353	AUGGUGGA CUGAUGAG X CGAA AUGGAUCU	179	AGATCCAT A TCCACCAT	591
1355	UAAUGGUG CUGAUGAG X CGAA AUAUGGAU	180	ATCCATAT C CACCATTA	592
1362	UGUCACAU CUGAUGAG X CGAA AUGGUGGA	181	TCCACCAT T ATGTGACA	593
1363	AUGUCACA CUGAUGAG X CGAA AAUGGUGG	182	CCACCATT A TGTGACAT	594
1372	GAUCCUUU CUGAUGAG X CGAA AUGUCACA	183	TGTGACAT T AAAGGATC	595

Table 9

1373	UGAUCCUU CUGAUGAG X CGAA AAUGUCAC	184	GTGACATT A AAGGATCA	596
1380	CUGUAUUAU CUGAUGAG X CGAA AUCCUUUA	185	TAAAGGAT C ATATACAG	597
1383	GCGCUGUA CUGAUGAG X CGAA AUGAUCCU	186	AGGATCAT A TACAGCGC	598
1385	UUGCGCUG CUGAUGAG X CGAA AUAUGAUC	187	GATCATAT A CAGCGCAA	599
1395	UAUGUUA CUGAUGAG X CGAA AUUGCGCU	188	AGCGCAAT T TGAACATA	600
1396	GUAUGUUC CUGAUGAG X CGAA AAUGCGC	189	GCGCAATT T GAACATAC	601
1403	CAGGAUGG CUGAUGAG X CGAA AUGUUCUA	190	TTGAACAT A CCATCCTG	602
1408	CGCAACAG CUGAUGAG X CGAA AUGGUAUG	191	CATACCAT C CTGTTGCG	603
1413	UUGGACGC CUGAUGAG X CGAA ACAGGAUG	192	CATCCTGT T GCGTCCAA	604
1418	ACAUGUUG CUGAUGAG X CGAA ACGCAACA	193	TGTTGCGT C CAACATGT	605
1427	AACUUCUU CUGAUGAG X CGAA ACAUGUUG	194	CAACATGT A AAGAAGTT	606
1435	CUGCUGAC CUGAUGAG X CGAA ACUUCUUU	195	AAAGAAGT T GTCAGCAG	607
1438	CCUCUGCU CUGAUGAG X CGAA ACAACUUC	196	GAAGTTGT C AGCAGAGG	608
1455	AAGUUUAA CUGAUGAG X CGAA AGUCAUCU	197	AGATGACT A TTAACTT	609
1457	CUAAGUUU CUGAUGAG X CGAA AUAGUCAU	198	ATGACTAT T AAACCTAG	610
1458	ACUAAGUU CUGAUGAG X CGAA AAUAGUCA	199	TGACTATT A AACTTAGT	611
1463	UUUGGACU CUGAUGAG X CGAA AGUUUAAU	200	ATTAACT T AGTCCAAA	612
1464	CUUUGGAC CUGAUGAG X CGAA AAGUUUAA	201	TTAACTT A GTCCAAAG	613
1467	UGGCUUUG CUGAUGAG X CGAA ACUAAGUU	202	AACTTAGT C CAAAGCCA	614
1479	AAGGUGUU CUGAUGAG X CGAA AGGUGGCU	203	AGCCACCT C AACACCTT	615
1487	AGAAAAUA CUGAUGAG X CGAA AGGUGUUG	204	CAACACCT T TATTTTCT	616
1488	CAGAAAAU CUGAUGAG X CGAA AAGGUGUU	205	AACACCTT T ATTTTCTG	617
1489	UCAGAAAA CUGAUGAG X CGAA AAAGGUGU	206	ACACCTTT A TTTTCTGA	618
1491	GCUCAGAA CUGAUGAG X CGAA AUAAAAGG	207	ACCTTTAT T TTCTGAGC	619
1492	AGCUCAGA CUGAUGAG X CGAA AAUAAAGG	208	CTTTTATT T TCTGAGCT	620
1493	AAGCUCAG CUGAUGAG X CGAA AAAUAAAG	209	CTTTATTT T CTGAGCTT	621
1494	AAAGCUCA CUGAUGAG X CGAA AAAAUAAA	210	TTTATTTT C TGAGCTTT	622
1501	UUCCAACA CUGAUGAG X CGAA AGCUCAGA	211	TCTGAGCT T TGTTGGAA	623
1502	UUUCCAAC CUGAUGAG X CGAA AAGCUCAG	212	CTGAGCTT T GTTGGAAA	624
1505	UGUUUUC CUGAUGAG X CGAA ACAAAGCU	213	AGCTTTGT T GGAAAACA	625
1518	AAUUCUGG CUGAUGAG X CGAA AUCAUGUU	214	AACATGAT A CCAGAATT	626
1526	GGCAAUU CUGAUGAG X CGAA AUUCUGGU	215	ACCAGAAT T AATTTGCC	627
1527	UGGCAAU CUGAUGAG X CGAA AAUUCUGG	216	CCAGAATT A ATTTGCCA	628
1530	AUGUGGCA CUGAUGAG X CGAA AUUAAUUC	217	GAATTAAT T TGCCACAT	629
1531	CAUGUGGC CUGAUGAG X CGAA AAUUAUUU	218	AATTAATT T GCCACATG	630
1541	AAACAGAC CUGAUGAG X CGAA ACAUGUGG	219	CCACATGT T GTCTGTTT	631
1544	UUAAAACA CUGAUGAG X CGAA ACAACAUG	220	CATGTTGT C TGTTTTAA	632
1548	ACUGUUAA CUGAUGAG X CGAA ACAGACAA	221	TTGTCTGT T TTAACAGT	633
1549	CACUGUUA CUGAUGAG X CGAA AACAGACA	222	TGTCTGTT T TAACAGTG	634
1550	CCACUGUU CUGAUGAG X CGAA AAACAGAC	223	GTCTGTTT T AACAGTGG	635
1551	UCCACUGU CUGAUGAG X CGAA AAAACAGA	224	TCTGTTTT A ACAGTGGA	636
1567	AAAAGUUA CUGAUGAG X CGAA ACAUGGGU	225	ACCCATGT A ATACTTTT	637
1570	GAUAAAAG CUGAUGAG X CGAA AUUACAUG	226	CATGTAAT A CTTTATC	638
1573	AUGGAUAA CUGAUGAG X CGAA AGUAUUAC	227	GTAATACT T TTATCCAT	639
1574	CAUGGAUA CUGAUGAG X CGAA AAGUAUUA	228	TAATACTT T TATCCATG	640
1575	ACAUGGAU CUGAUGAG X CGAA AAAGUAUU	229	AATACTTT T ATCCATGT	641
1576	AACAUGGA CUGAUGAG X CGAA AAAAGUAU	230	ATACTTTT A TCCATGTT	642

Table 9

1578	UAAACAUG CUGAUGAG X CGAA AUAAAAGU	231	ACTTTTAT C CATGTTTA	643
1584	CUUUUUUA CUGAUGAG X CGAA ACAUGGAU	232	ATCCATGT T TAAAAAAG	644
1585	UCUUUUUU CUGAUGAG X CGAA AACAUUGA	233	TCCATGTT T AAAAAAGA	645
1586	UUCUUUUU CUGAUGAG X CGAA AAACAUGG	234	CCATGTTT A AAAAAGAA	646
1600	UUUGUCCA CUGAUGAG X CGAA AUUCCUUC	235	GAAGGAAT T TGGACAAA	647
1601	CUUUGUCC CUGAUGAG X CGAA AAUCCUUC	236	AAGGAATT T GGACAAAG	648
1619	UUACAUAU CUGAUGAG X CGAA ACGGUUUG	237	CAAACCGT C TAATGTAA	649
1621	AAUUAUUA CUGAUGAG X CGAA AGACGGUU	238	AACCGTCT A ATGTAATT	650
1626	UGGUUAAU CUGAUGAG X CGAA ACAUUAGA	239	TCTAATGT A ATTAACCA	651
1629	CGUUGGUU CUGAUGAG X CGAA AUUACAUU	240	AATGTAAT T AACCAACG	652
1630	UCGUUGGU CUGAUGAG X CGAA AAUUAUUA	241	ATGTAATT A ACCAACGA	653
1646	AGUCCGGA CUGAUGAG X CGAA AGCUUUUU	242	AAAAAGCT T TCCGACT	654
1647	AAGUCCGG CUGAUGAG X CGAA AAGCUUUU	243	AAAAGCTT T CCGGACTT	655
1648	AAAGUCCG CUGAUGAG X CGAA AAAGCUUU	244	AAAGCTTT C CGGACTTT	656
1655	GCAUUUAA CUGAUGAG X CGAA AGUCCGGA	245	TCCGACT T TTAATGCT	657
1656	AGCAUUUA CUGAUGAG X CGAA AAGUCCGG	246	CCGGACTT T TAAATGCT	658
1657	UAGCAUUU CUGAUGAG X CGAA AAAGUCCG	247	CGGACTTT T AAATGCTA	659
1658	UUAGCAUU CUGAUGAG X CGAA AAAAGUCC	248	GGACTTTT A AATGCTAA	660
1665	AAAACAGU CUGAUGAG X CGAA AGCAUUUA	249	TAAATGCT A ACTGTTTT	661
1671	AGGGGAAA CUGAUGAG X CGAA ACAGUUAG	250	CTAACTGT T TTTCCCT	662
1672	AAGGGGAA CUGAUGAG X CGAA AACAGUUA	251	TAAGTGT T TTTCCCTT	663
1673	GAAGGGGA CUGAUGAG X CGAA AAACAGUU	252	AACTGTTT T TCCCTTC	664
1674	GGAAGGGG CUGAUGAG X CGAA AAAACAGU	253	ACTGTTTT T CCCCTTCC	665
1675	AGGAAGGG CUGAUGAG X CGAA AAAACAG	254	CTGTTTTT C CCCTTCT	666
1680	UAGACAGG CUGAUGAG X CGAA AGGGGAAA	255	TTTCCCTT T CCTGTCTA	667
1681	CUAGACAG CUGAUGAG X CGAA AAGGGGAA	256	TTCCCTT C CTGTCTAG	668
1686	UUUUCUA CUGAUGAG X CGAA ACAGGAAG	257	CTTCTGT C TAGGAAAA	669
1688	CAUUUUC CUGAUGAG X CGAA AGACAGGA	258	TCCTGTCT A GGAAAAATG	670
1699	GAGCUUUA CUGAUGAG X CGAA AGCAUUUU	259	AAAATGCT A TAAAGCTC	671
1701	UUGAGCUU CUGAUGAG X CGAA AUAGCAUU	260	AATGCTAT A AAGCTCAA	672
1707	ACUAAUUU CUGAUGAG X CGAA AGCUUUUU	261	ATAAAGCT C AAATTAGT	673
1712	UCCUAAU CUGAUGAG X CGAA AUUUGAGC	262	GCTCAAAT T AGTTAGGA	674
1713	UUCCUAA CUGAUGAG X CGAA AAUUGAG	263	CTCAAAT A GTTAGGAA	675
1716	UCAUUCU CUGAUGAG X CGAA ACUAAUUU	264	AAATTAGT T AGGAATGA	676
1717	GUCAUUC CUGAUGAG X CGAA AACUAAUU	265	AATTAGTT A GGAATGAC	677
1727	AAACGUUU CUGAUGAG X CGAA AGUCAUUC	266	GAATGACT T ATACGTTT	678
1728	AAAACGUA CUGAUGAG X CGAA AAGUCAUU	267	AATGACTT A TACGTTTT	679
1730	ACAAAACG CUGAUGAG X CGAA AUAAGUCA	268	TGACTTAT A CGTTTTGT	680
1734	CAAAAACA CUGAUGAG X CGAA ACGUAUUA	269	TTATACGT T TGTTTTGT	681
1735	UCAAAAAC CUGAUGAG X CGAA AACGUUAU	270	TATACGTT T TGTTTTGA	682
1736	UUCAAAAC CUGAUGAG X CGAA AAACGUUU	271	ATACGTTT T GTTTTGAA	683
1739	GUUUUCAA CUGAUGAG X CGAA ACAAACG	272	CGTTTTGT T TGAATAC	684
1740	GGUUAUUA CUGAUGAG X CGAA AACAAAAC	273	GTTTTGTT T TGAATACC	685
1741	AGGUUUUC CUGAUGAG X CGAA AAACAAAA	274	TTTTGTTT T GAATACCT	686
1746	CUCUUAGG CUGAUGAG X CGAA AUUCAAAA	275	TTTTGAAT A CCTAAGAG	687
1750	GUUUCUCU CUGAUGAG X CGAA AGGUUUUC	276	GAATACCT A AGAGATAC	688
1757	CCAAAAG CUGAUGAG X CGAA AUCUUUA	277	TAAGAGAT A CTTTTTGG	689

Table 9

1760	UAUCCAAA CUGAUGAG X CGAA AGUAUCUC	278	GAGATACT T TTTGGATA	690
1761	AUAUCCAA CUGAUGAG X CGAA AAGUAUCU	279	AGATACTT T TTGGATAT	691
1762	AAUAUCCA CUGAUGAG X CGAA AAAGUAUC	280	GATACTTT T TGGATATT	692
1763	AAAUAUCC CUGAUGAG X CGAA AAAAGUAU	281	ATACTTTT T GGATATTT	693
1768	AAUAUAAA CUGAUGAG X CGAA AUCCAAAA	282	TTTGGAT A TTTATATT	694
1770	GCAAUUAU CUGAUGAG X CGAA AUAUCCAA	283	TTGGATAT T TATATGTC	695
1771	GGCAAUUAU CUGAUGAG X CGAA AAUAUCCA	284	TGGATATT T ATATTGCC	696
1772	UGGCAUAU CUGAUGAG X CGAA AAAUAUCC	285	GGATATTT A TATTGCCA	697
1774	UAUGGCAA CUGAUGAG X CGAA AUAAAUAU	286	ATATTTAT A TTGCCATA	698
1776	AAUAUGGC CUGAUGAG X CGAA AUUAUAAU	287	ATTTATAT T GCCATATT	699
1782	AGUAAGAA CUGAUGAG X CGAA AUGGCAU	288	ATTGCCAT A TTCTTACT	700
1784	CAAGUAAG CUGAUGAG X CGAA AU AUGGCA	289	TGCCATAT T CTTACTTG	701
1785	UCAAGUAA CUGAUGAG X CGAA AAUAUGGC	290	GCCATATT C TTACTTGA	702
1787	AUUCAAGU CUGAUGAG X CGAA AGAAUAUG	291	CATATTCT T ACTTGAAT	703
1788	CAUUCAAG CUGAUGAG X CGAA AAGAAUAU	292	ATATTCTT A CTTGAATG	704
1791	AAGCAUUC CUGAUGAG X CGAA AGUAAGAA	293	TTCTTACT T GAATGCTT	705
1799	GUCAUUCA CUGAUGAG X CGAA AGCAUUCA	294	TGAATGCT T TGAATGAC	706
1800	AGUCAUUC CUGAUGAG X CGAA AAGCAUUC	295	GAATGCTT T GAATGACT	707
1809	ACUGGAUG CUGAUGAG X CGAA AGUCAUUC	296	GAATGACT A CATCCAGT	708
1813	CAGAACUG CUGAUGAG X CGAA AUGUAGUC	297	GACTACAT C CAGTTCTG	709
1818	AGGUGCAG CUGAUGAG X CGAA ACUGGAUG	298	CATCCAGT T CTGCACCT	710
1819	UAGGUGCA CUGAUGAG X CGAA AACUGGAU	299	ATCCAGTT C TGCACCTA	711
1827	AGAGGGUA CUGAUGAG X CGAA AGGUGCAG	300	CTGCACCT A TACCCTCT	712
1829	CCAGAGGG CUGAUGAG X CGAA AUAGGUGC	301	GCACCTAT A CCCTCTGG	713
1834	CAACACCA CUGAUGAG X CGAA AGGUUAUA	302	TATACCTT C TGGTGTG	714
1841	UAAAAAGC CUGAUGAG X CGAA ACACCAGA	303	TCTGGTGT T GCTTTTFA	715
1845	AGGUUAAA CUGAUGAG X CGAA AGCAACAC	304	GTGTTGCT T TTAACCT	716
1846	AAGGUUAA CUGAUGAG X CGAA AAGCAACA	305	TGTTGCTT T TTAACCTT	717
1847	GAAGGUUA CUGAUGAG X CGAA AAAGCAAC	306	GTTGCTTT T TAACCTTC	718
1848	GGAAGGUU CUGAUGAG X CGAA AAAAGCAA	307	TTGCTTTT T AACCTTCC	719
1849	AGGAAGGU CUGAUGAG X CGAA AAAAGCA	308	TGCTTTT A ACCTTCT	720
1854	AUUC CAGG CUGAUGAG X CGAA AGGUUAAA	309	TTTAACCT T CCTGGAAT	721
1855	GAUUC CAG CUGAUGAG X CGAA AAGGUUAA	310	TTAACCTT C CTGGAATC	722
1863	AGAAA AUG CUGAUGAG X CGAA AUUC CAGG	311	CCTGGAAT C CATTTTCT	723
1867	UUUUAGAA CUGAUGAG X CGAA AUGGAUUC	312	GAATCCAT T TTCTAAAA	724
1868	UUUUUAGA CUGAUGAG X CGAA AAUGGAUU	313	AATCCATT T TCTAAAAA	725
1869	UUUUUAG CUGAUGAG X CGAA AAAUGGAU	314	ATCCATT T CTAAAAAA	726
1870	AUUUUUA CUGAUGAG X CGAA AAAAUGGA	315	TCCATTT C TAAAAAAT	727
1872	UUUUUUU CUGAUGAG X CGAA AGAAAAUG	316	CATTTTCT A AAAAATAA	728
1879	UGUGUCUU CUGAUGAG X CGAA AUUUUUUA	317	TAAAAAAT A AAGACACA	729
1889	CUGAGAAG CUGAUGAG X CGAA AUGUGUCU	318	AGACACAT T CTCTCAG	730
1890	GCUGAGAA CUGAUGAG X CGAA AAUGUGUC	319	GACACATT C TTCTCAGC	731
1892	GUGCUGAG CUGAUGAG X CGAA AGAAUGUG	320	CACATTCT T CTCAGCAC	732
1893	GGUGCUGA CUGAUGAG X CGAA AAGAAUGU	321	ACATTCTT C TCAGCACC	733
1895	GUGGUGCU CUGAUGAG X CGAA AGAAGAAU	322	ATTCTTCT C AGCACCAC	734
1913	UUUUGGAA CUGAUGAG X CGAA AGGUGUUG	323	CAACACCT A TTCCAAAA	735
1915	GAUUUUGG CUGAUGAG X CGAA AUAGGUGU	324	ACACCTAT T CCAAAATC	736

Table 9

1916	CGAUUUUG CUGAUGAG X CGAA AAUAGGUG	325	CACCTATT C CAAAATCG	737
1923	AUGUGGUC CUGAUGAG X CGAA AUUUUGGA	326	TCCAAAAT C GACCACAT	738
1932	CUUCCAAA CUGAUGAG X CGAA AUGUGGUC	327	GACCACAT A TTTGGAAG	739
1934	UACUCCAA CUGAUGAG X CGAA AUAUGUGG	328	CCACATAT T TGGAAGTA	740
1935	UUACUCC CUGAUGAG X CGAA AAUAUGUG	329	CACATATT T GGAAGTAA	741
1942	GAGAGCUU CUGAUGAG X CGAA ACUUCCAA	330	TTGGAAGT A AAGCTCTC	742
1948	GCUGAGGA CUGAUGAG X CGAA AGCUUUAC	331	GTAAGCT C TCCTCAGC	743
1950	UUGCUGAG CUGAUGAG X CGAA AGAGCUUU	332	AAAGCTCT C CTCAGCAA	744
1953	CAUUGUCU CUGAUGAG X CGAA AGGAGAGC	333	GCTCTCCT C AGCAAATG	745
1963	UGUUCUUU CUGAUGAG X CGAA ACAUUUGC	334	GCAAATGT A AAAGAACA	746
1977	UUUGUUAU CUGAUGAG X CGAA AUUUCUGU	335	ACAGAAAT T ATAACAAA	747
1978	GUUUGUUA CUGAUGAG X CGAA AAUUCUG	336	CAGAAATT A TAACAAAC	748
1980	CAGUUUGU CUGAUGAG X CGAA AUAAUUUC	337	GAAATTAT A ACAAACTG	749
1990	GUCUGAGA CUGAUGAG X CGAA ACAGUUUG	338	CAAACTGT C TCTCAGAC	750
1992	UGGUCUGA CUGAUGAG X CGAA AGACAGUU	339	AACTGTCT C TCAGACCA	751
1994	UGUGGUCU CUGAUGAG X CGAA AGAGACAG	340	CTGTCTCT C AGACCACA	752
2005	UUUGGUUA CUGAUGAG X CGAA ACUGUGGU	341	ACCACAGT A TAACCAAA	753
2007	AGUUUGGU CUGAUGAG X CGAA AUACUGUG	342	CACAGTAT A ACCAAACT	754
2016	CUGAGUUC CUGAUGAG X CGAA AGUUUGGU	343	ACCAAACT A GAACTCAG	755
2022	UUAUCCU CUGAUGAG X CGAA AGUUCUAG	344	CTAGAACT C AGGATTAA	756
2028	AGUUUCUU CUGAUGAG X CGAA AUCCUGAG	345	CTCAGGAT T AAGAACT	757
2029	GAGUUUCU CUGAUGAG X CGAA AAUCCUGA	346	TCAGGATT A AGAACTC	758
2037	UUUUGAGU CUGAUGAG X CGAA AGUUUCUU	347	AAGAACT C ACTCAAAA	759
2041	GUGGUUUU CUGAUGAG X CGAA AGUGAGUU	348	AACTCACT C AAAACCAC	760
2056	UUUCCAUG CUGAUGAG X CGAA AGUUGUGU	349	ACACAACT A CATGGAAA	761
2079	UCAUUCAG CUGAUGAG X CGAA AGCAGGUU	350	AACCTGCT C CTGAATGA	762
2090	GUAUCCAG CUGAUGAG X CGAA AGUCAUUC	351	GAATGACT A CTGGATAC	763
2097	UUGUUAUG CUGAUGAG X CGAA AUCCAGUA	352	TACTGGAT A CATAACAA	764
2101	CAUUUUGU CUGAUGAG X CGAA AUGUAUCC	353	GGATACAT A ACAAAATG	765
2121	AAAUUCUU CUGAUGAG X CGAA AUUUCUGC	354	GCAGAAAT A AAGATGTT	766
2129	UUUUAAG CUGAUGAG X CGAA ACAUCUUU	355	AAAGATGT T CTTAAAA	767
2130	GUUUUAAA CUGAUGAG X CGAA AACAUCUU	356	AAGATGTT C TTTAAAC	768
2132	UGGUUUUA CUGAUGAG X CGAA AGAACAUC	357	GATGTTCT T TAAACCA	769
2133	UUGGUUUU CUGAUGAG X CGAA AAGAACAU	358	ATGTTCTT T AAAACCAA	770
2134	AUUGGUUU CUGAUGAG X CGAA AAAGAACA	359	TGTTCTTT A AAACCAAT	771
2162	GAUUCUGG CUGAUGAG X CGAA AUGUUGUG	360	CACAACAT A CCAGAATC	772
2170	GUCCCAGA CUGAUGAG X CGAA AUUCUGGU	361	ACCAGAAT C TCTGGGAC	773
2172	GUGUCCCA CUGAUGAG X CGAA AGAUUCUG	362	CAGAATCT C TGGGACAC	774
2183	CUGCUUUG CUGAUGAG X CGAA AUGUGUCC	363	GGACACAT T CAAAGCAG	775
2184	ACUGCUUU CUGAUGAG X CGAA AAUGUGUC	364	GACACATT C AAAGCAGT	776
2197	UUUCCUC CUGAUGAG X CGAA ACACACUG	365	CAGTGTGT A GAGGGAAA	777
2207	GUGCUAUA CUGAUGAG X CGAA AUUUCCCU	366	AGGGAAAT T TATAGCAC	778
2208	AGUGCUAU CUGAUGAG X CGAA AAUUUCCC	367	GGGAAATT T ATAGCACT	779
2209	UAGUGCUA CUGAUGAG X CGAA AAAUUUCC	368	GGAAATTT A TAGCACTA	780
2211	UUUAGUGC CUGAUGAG X CGAA AUAAAUUU	369	AAATTTAT A GCACTAAA	781
2217	UGGGCAUU CUGAUGAG X CGAA AGUGCUAU	370	ATAGCACT A AATGCCCA	782
2244	AUUUJAGA CUGAUGAG X CGAA AUUUCCUG	371	CAGGAAAT A TCTAAAT	783

Table 9

2246	CAAUUUUA CUGAUGAG X CGAA AUAUUUCC	372	GGAAATAT C TAAAATTG	784
2248	GUCAAUUU CUGAUGAG X CGAA AGAUAUUU	373	AAATATCT A AAATTGAC	785
2253	AGGGUGUC CUGAUGAG X CGAA AUUUUAGA	374	TCTAAAAT T GACACCCT	786
2262	UGUGAUGU CUGAUGAG X CGAA AGGGUGUC	375	GACACCCT A ACATCACA	787
2267	UUAUUUGU CUGAUGAG X CGAA AUGUUAGG	376	CCTAACAT C ACAATTAA	788
2273	GUUCUUUU CUGAUGAG X CGAA AUUGUGAU	377	ATCACAAT T AAAAGAAC	789
2274	AGUUCUUU CUGAUGAG X CGAA AAUUGUGA	378	TCACAATT A AAAGAACT	790
2283	UGCUUCUC CUGAUGAG X CGAA AGUUCUUU	379	AAAGAACT A GAGAAGCA	791
2305	AGCUUUUC CUGAUGAG X CGAA AUGUGUUU	380	AAACACAT T GAAAAGCT	792
2314	CCUUCUCU CUGAUGAG X CGAA AGCUUUUC	381	GAAAAGCT A AGAGAAGG	793
2331	AUCUUAGU CUGAUGAG X CGAA AUUUCUUG	382	CAAGAAAT A ACTAAGAT	794
2335	UCUGAUCU CUGAUGAG X CGAA AGUUUUUU	383	AAATAACT A AGATCAGA	795
2340	UCUGCUCU CUGAUGAG X CGAA AUCUUAGU	384	ACTAAGAT C AGAGCAGA	796
2361	UGUGUCUC CUGAUGAG X CGAA AUUCCUUU	385	AAGGAAAT A GAGACACA	797
2377	UUUUUGAA CUGAUGAG X CGAA AGUUUUUU	386	AAAAAACT C TTCAAAAA	798
2379	AUUUUUUG CUGAUGAG X CGAA AGAGUUUU	387	AAAACTCT T CAAAAAAT	799
2380	GAUUUUUU CUGAUGAG X CGAA AAGAGUUU	388	AAACTCTT C AAAAAATC	800
2388	GAUUCAUU CUGAUGAG X CGAA AUUUUUUG	389	CAAAAAAT C AATGAATC	801
2396	AGCUCCUG CUGAUGAG X CGAA AUUCAUUG	390	CAATGAAT C CAGGAGCT	802
2408	UUUCAAAA CUGAUGAG X CGAA ACCAGCUC	391	GAGCTGGT T TTTTGAAA	803
2409	GUUUCAAA CUGAUGAG X CGAA AACCAGCU	392	AGCTGGTT T TTTGAAAC	804
2410	CGUUUCAA CUGAUGAG X CGAA AAACCAGC	393	GCTGGTTT T TTGAAACG	805
2411	UCGUUUCA CUGAUGAG X CGAA AAAACCAG	394	CTGGTTTT T TGAAACGA	806
2412	AUCGUUUC CUGAUGAG X CGAA AAAAACCA	395	TGGTTTTT T GAAACGAT	807
2421	AUUUUGUU CUGAUGAG X CGAA AUCGUUUC	396	GAAACGAT C AACAAAAT	808
2430	UGUCUAUC CUGAUGAG X CGAA AUUUUGUU	397	AACAAAAT T GATAGACA	809
2434	CUAGUGUC CUGAUGAG X CGAA AUCAAUUU	398	AAATTGAT A GACACTAG	810
2441	AGUCUUGC CUGAUGAG X CGAA AGUGUCUA	399	TAGACACT A GCAAGACT	811
2450	UUCUUUAU CUGAUGAG X CGAA AGUCUUGC	400	GCAAGACT A ATAAAGAA	812
2453	UUCUUCUU CUGAUGAG X CGAA AUUAGUCU	401	AGACTAAT A AAGAAGAA	813
2475	UUCUAUUU CUGAUGAG X CGAA AUUCUUCU	402	AGAAGAAT C AAATAGAA	814
2480	AUUGCUUC CUGAUGAG X CGAA AUUUGAUU	403	AATCAAAT A GAAGCAAT	815
2489	UCAUUUUU CUGAUGAG X CGAA AUUGCUUC	404	GAAGCAAT A AAAATGA	816
2499	AUCCCCUU CUGAUGAG X CGAA AUCAUUUU	405	AAAATGAT A AAGGGGAT	817
2508	GGUGGUGA CUGAUGAG X CGAA AUCCCCUU	406	AAGGGGAT A TCACCACC	818
2510	UUGGUGGU CUGAUGAG X CGAA AUAUCCCC	407	GGGGATAT C ACCACCAA	819
2520	UUCUGUGG CUGAUGAG X CGAA AUUGGUGG	408	CCACCAAT C CCACAGAA	820
2531	UGGUGGUU CUGAUGAG X CGAA AUUUCUGU	409	ACAGAAAT A AACCACCA	821
2541	UAUUCUCU CUGAUGAG X CGAA AUGGUGGU	410	ACCACCAT C AGAGAATA	822
2549	GUUUGUAG CUGAUGAG X CGAA AUUCUCUG	411	CAGAGAAT A CTACAAAC	823
2552	GGUGUUUG CUGAUGAG X CGAA AGUAUUCU	412	AGAATACT A CAAACACC	824

Input Sequence = HSU29607. Cut Site = UH/.

Stem Length = 8. Core Sequence = CUGAUGAG X CGAA (X = GCCGUUAGGC or other stem II)

Seq1 = HSU29607 (Human methionine aminopeptidase mRNA, complete cds., 2569 bp)

Table 10

Table 10: Human methionine aminopeptidase type 2 (MetAP-2) NCH Ribozyme and Target Sequence

Nt. position	Ribozyme Sequence	Seq. ID Nos.	Substrate Sequence	Seq. ID Nos.
10	CCCGAGAG CUGAUGAG X CGAA IACGAGGG	825	CCCTCGTC T CTCTCGGG	1255
12	UGCCCGAG CUGAUGAG X CGAA IAGACGAG	826	CTCGTCTC T CTCGGGCA	1256
14	GUUGCCCG CUGAUGAG X CGAA IAGAGACG	827	CGTCTCTC T CCGGCAAC	1257
20	CGCCAUGU CUGAUGAG X CGAA ICCCGAGA	828	TCTCGGGC A ACATGGCG	1258
23	GCCCGCCA CUGAUGAG X CGAA IUUGCCCG	829	CGGGCAAC A TGGCGGGC	1259
49	CUCCCGGA CUGAUGAG X CGAA ICCGCUAC	830	GTAGCGGC C TCGGGAG	1260
50	GCUCCCGG CUGAUGAG X CGAA IGC CGCUA	831	TAGCGGCC T CCGGGAGC	1261
52	UGGCUCCC CUGAUGAG X CGAA IAGGCCGC	832	GCGGCCTC C GGGAGCCA	1262
59	AUUCAGGU CUGAUGAG X CGAA ICUCCCGG	833	CCGGGAGC C ACCTGAAT	1263
60	CAUUCAGG CUGAUGAG X CGAA IGCUCCCG	834	CGGGAGCC A CCTGAATG	1264
62	GCCAUUCA CUGAUGAG X CGAA IUUGCUCC	835	GGAGCCAC C TGAATGGC	1265
63	CGCCAUUC CUGAUGAG X CGAA IGUGGCUC	836	GAGCCACC T GAATGGCG	1266
74	UGGAUCCA CUGAUGAG X CGAA IUCGCAU	837	ATGGCGAC C TGATCCA	1267
75	CUGGAUCC CUGAUGAG X CGAA IGUCGCCA	838	TGGCGACC T GGATCCAG	1268
81	UGUCGUCU CUGAUGAG X CGAA IAUCCAGG	839	CCTGGATC C AGACGACA	1269
82	CUGUCGUC CUGAUGAG X CGAA IGAUCCAG	840	CTGGATCC A GACGACAG	1270
89	UUCUCCCC CUGAUGAG X CGAA IUCGUCUG	841	CAGACGAC A GGAAGAA	1271
103	GUAGAGGC CUGAUGAG X CGAA ICUCCUUC	842	GAAGGAGC T GCCTCTAC	1272
106	GCCGUAGA CUGAUGAG X CGAA ICAGCUCC	843	GGAGCTGC C TCTACGGC	1273
107	AGCCGUAG CUGAUGAG X CGAA IGCAGCUC	844	GAGCTGCC T CTACGGCT	1274
109	UCAGCCGU CUGAUGAG X CGAA IAGGCAGC	845	GCTGCCTC T ACGGCTGA	1275
115	GCUUCCUC CUGAUGAG X CGAA ICCGUAGA	846	TCTACGGC T GAGGAAGC	1276
124	UUCUUGGC CUGAUGAG X CGAA ICUUCCUC	847	GAGGAAGC A GCCAAGAA	1277
127	UUUUUCUU CUGAUGAG X CGAA ICUGCUUC	848	GAAGCAGC C AAGAAAAA	1278
128	UUUUUCUU CUGAUGAG X CGAA IGCUGCUU	849	AAGCAGCC A AGAAAAA	1279
158	AGGCCUUU CUGAUGAG X CGAA ICUCUUCU	850	AGAAGAGC A AAGGGCCT	1280
165	CUGCAGAA CUGAUGAG X CGAA ICCUUUUG	851	CAAAGGGC C TTCTGCAG	1281
166	GCUGCAGA CUGAUGAG X CGAA IGCCUUU	852	AAAGGGCC T TCTGCAGC	1282
169	CCUGCUGC CUGAUGAG X CGAA IAAGGCC	853	GGGCCTTC T GCAGCAGG	1283
172	UCCCCUGC CUGAUGAG X CGAA ICAGAAGG	854	CCTTCTGC A GCAGGGGA	1284
175	UGUCCCC CUGAUGAG X CGAA ICUGCAGA	855	TCTGCAGC A GGGGAACA	1285
183	CAGGUUCC CUGAUGAG X CGAA IUUCCCCU	856	AGGGGAAC A GGAACCTG	1286
189	CUUUAUCA CUGAUGAG X CGAA IUUCCUGU	857	ACAGGAAC C TGATAAAG	1287
190	UCUUUAUC CUGAUGAG X CGAA IGUCCUG	858	CAGGAACC T GATAAAGA	1288
202	GAGGCUCC CUGAUGAG X CGAA IAUUCUUU	859	AAAGAATC A GGAGCCTC	1289
208	UCCACUGA CUGAUGAG X CGAA ICUCCUGA	860	TCAGGAGC C TCAGTGGA	1290
209	AUCCACUG CUGAUGAG X CGAA IGCUCUG	861	CAGGAGCC T CAGTGGAT	1291
211	UCAUCCAC CUGAUGAG X CGAA IAGGCUCC	862	GGAGCCTC A GTGGATGA	1292
226	AACUGUCU CUGAUGAG X CGAA ICUACUUC	863	GAAGTAGC A AGACAGTT	1293
231	UUUCCAAC CUGAUGAG X CGAA IUUCUUCU	864	AGCAAGAC A GTTGGAAA	1294
244	UCCAAUGC CUGAUGAG X CGAA IAUUCUUC	865	GAAAGATC A GCATTGGA	1295
247	UCUCCAA CUGAUGAG X CGAA ICUGAUCU	866	AGATCAGC A TTGGAAGA	1296

Table 10

307	UUUCCAGU CUGAUGAG X CGAA ICUCCAUC	867	GATGGAGC A ACTGGAAA	1297
310	UUCUUUCC CUGAUGAG X CGAA IUUGCUC	868	GGAGCAAC T GGAAAGAA	1298
348	GAACUUUU CUGAUGAG X CGAA IUCCUCUC	869	GAGAGGAC C AAAAGTTC	1299
349	UGAACUUU CUGAUGAG X CGAA IGUCCUCU	870	AGAGGACC A AAAGTCA	1300
357	GGUCUGUU CUGAUGAG X CGAA IAAUUUU	871	AAAAGTTC A AACAGACC	1301
361	GGAGGGUC CUGAUGAG X CGAA IUUUGAAC	872	GTTCAAAC A GACCCTCC	1302
365	UGAGGGAG CUGAUGAG X CGAA IUCUGUUU	873	AAACAGAC C CTCCTCA	1303
366	CUGAGGGA CUGAUGAG X CGAA IGUCUGUU	874	AACAGACC C TCCCTCAG	1304
367	ACUGAGGG CUGAUGAG X CGAA IGGUCUGU	875	ACAGACCC T CCCTCAGT	1305
369	GAACUGAG CUGAUGAG X CGAA IAGGGUCU	876	AGACCCTC C CTCAGTTC	1306
370	GGAACUGA CUGAUGAG X CGAA IGAGGGUC	877	GACCCTCC C TCAGTTCC	1307
371	UGGAACUG CUGAUGAG X CGAA IGGAGGGU	878	ACCCTCCC T CAGTTCCA	1308
373	AUUGGAAC CUGAUGAG X CGAA IAGGGAGG	879	CCTCCCTC A GTTCCAAT	1309
378	CACAUUUU CUGAUGAG X CGAA IAAUCUGAG	880	CTCAGTTC C AATATGTG	1310
379	UCACAUUU CUGAUGAG X CGAA IGAACUGA	881	TCAGTTCC A ATATGTGA	1311
389	AGGAUACA CUGAUGAG X CGAA IUCACAUU	882	TATGTGAC C TGTATCCT	1312
390	UAGGAUAC CUGAUGAG X CGAA IGUCACAU	883	ATGTGACC T GTATCCTA	1313
396	CACCAUUA CUGAUGAG X CGAA IAUACAGG	884	CCTGTATC C TAATGGTG	1314
397	ACACCAUU CUGAUGAG X CGAA IGAUACAG	885	CTGTATCC T AATGGTGT	1315
411	GUCCUUUG CUGAUGAG X CGAA IAAUACA	886	TGTATTTC C CAAAGGAC	1316
412	UGUCCUUU CUGAUGAG X CGAA IGAAAUAC	887	GTATTTCC C AAAGGACA	1317
413	UUGUCCUU CUGAUGAG X CGAA IGGAAUA	888	TATTTCCC A AAGGACAA	1318
420	CGCAUUCU CUGAUGAG X CGAA IUCCUUUG	889	CAAAGGAC A AGAATGCG	1319
434	UGUGGGUG CUGAUGAG X CGAA IUUAUUCG	890	GCGAATAC C CACCCACA	1320
435	GUGUGGGU CUGAUGAG X CGAA IGUAUUCG	891	CGAATACC C ACCCACAC	1321
436	UGUGUGGG CUGAUGAG X CGAA IGGUAUUC	892	GAATACCC A CCCACACA	1322
438	CUUGUGUG CUGAUGAG X CGAA IUGGGUUA	893	ATACCCAC C CACACAAG	1323
439	UCUUGUGU CUGAUGAG X CGAA IGUGGGUA	894	TACCCACC C ACACAAGA	1324
440	AUCUUGUG CUGAUGAG X CGAA IGGUGGGU	895	ACCCACCC A CACAAGAT	1325
442	CCAUCUUG CUGAUGAG X CGAA IUGGGUGG	896	CCACCCAC A CAAGATGG	1326
444	GCCCAUCU CUGAUGAG X CGAA IUGUGGGU	897	ACCCACAC A AGATGGGC	1327
457	CAAGCAGC CUGAUGAG X CGAA IUUCGCCC	898	GGGCGAAC A GCTGCTTG	1328
460	CUCCAAGC CUGAUGAG X CGAA ICUGUUCG	899	CGAACAGC T GCTTGGAG	1329
463	GUUCUCCA CUGAUGAG X CGAA ICAGCUGU	900	ACAGCTGC T TGGAGAAC	1330
472	UCACUUGU CUGAUGAG X CGAA IUUCUCCA	901	TGGAGAAC T ACAAGTGA	1331
475	UCUUCACU CUGAUGAG X CGAA IUAGUUCU	902	AGAACTAC A AGTGAAGA	1332
493	UGAUCUAA CUGAUGAG X CGAA ICUUUCUU	903	AAGAAAGC A TTAGATCA	1333
501	CACUUGCC CUGAUGAG X CGAA IAUCUAAU	904	ATTAGATC A GGCAAGTG	1334
505	UCUUCACU CUGAUGAG X CGAA ICCUGAUC	905	GATCAGGC A AGTGAAGA	1335
538	GCUUCUGC CUGAUGAG X CGAA ICUUCUCG	906	CGAGAAGC T GCAGAAGC	1336
541	UGUGCUUC CUGAUGAG X CGAA ICAGCUUC	907	GAAGCTGC A GAAGCACA	1337
547	UGUCGAUG CUGAUGAG X CGAA ICUUCUGC	908	GCAGAAGC A CATCGACA	1338
549	CUUGUCGA CUGAUGAG X CGAA IUGCUUCU	909	AGAAGCAC A TCGACAAG	1339
555	UUCUAACU CUGAUGAG X CGAA IUCGAUGU	910	ACATCGAC A AGTTAGAA	1340
578	CUUGAUCC CUGAUGAG X CGAA ICUCAUUA	911	TAATGAGC T GGATCAAG	1341
584	CCCAGGCU CUGAUGAG X CGAA IAUCCAGC	912	GCTGGATC A AGCCTGGG	1342
588	UCAUCCCA CUGAUGAG X CGAA ICUUGAUC	913	GATCAAGC C TGGGATGA	1343

Table 10

589	GUCAUCCC CUGAUGAG X CGAA IGCUUGAU	914	ATCAAGCC T GGGATGAC	1344
598	UCUAUCAU CUGAUGAG X CGAA IUCAUCCC	915	GGGATGAC A ATGATAGA	1345
611	CUUUUCAC CUGAUGAG X CGAA IAUUUCUA	916	TAGAAATC T GTGAAAAG	1346
629	GCGUGAAC CUGAUGAG X CGAA IUCUUCCA	917	TGGAAGAC T GTTCACGC	1347
634	AACUUGCG CUGAUGAG X CGAA IAACAGUC	918	GACTGTTC A CGCAAGTT	1348
638	UAUUAACU CUGAUGAG X CGAA ICGUGAAC	919	GTTACACG A AGTTAATA	1349
667	GCCAGGCC CUGAUGAG X CGAA ICAUUUAA	920	TTAAATGC A GGCCTGGC	1350
671	AAAUGCCA CUGAUGAG X CGAA ICCUGCAU	921	ATGCAGGC C TGGCATT	1351
672	GAAUGGCC CUGAUGAG X CGAA IGCCUGCA	922	TGCAGGCC T GGCATTTC	1352
676	GUAGGAAA CUGAUGAG X CGAA ICCAGGCC	923	GGCCTGGC A TTTCTAC	1353
681	AUCCAGUA CUGAUGAG X CGAA IAAAUGCC	924	GGCATTTC C TACTGGAT	1354
682	CAUCCAGU CUGAUGAG X CGAA IGAAUUGC	925	GCATTTC C T ACTGGATG	1355
685	GAACAUC CUGAUGAG X CGAA IUAGGAAA	926	TTTCCTAC T GGATGTT	1356
694	UUAUUGAG CUGAUGAG X CGAA IAACAUC	927	GGATGTT T CTCAATAA	1357
696	AAUUAUUG CUGAUGAG X CGAA IAGAACA	928	ATGTTCTC T CAATAATT	1358
698	ACAUAUUA CUGAUGAG X CGAA IAGAGAAC	929	GTTCTCTC A ATAATTGT	1359
709	UAAUGGGC CUGAUGAG X CGAA ICACAAUU	930	AATTGTGC T GCCCATT	1360
712	GUUAUAUG CUGAUGAG X CGAA ICAGCACA	931	TGTGCTGC C CATTATAC	1361
713	AGUAUAAU CUGAUGAG X CGAA IGCAGCAC	932	GTGCTGCC C ATTATACT	1362
714	GAGUAUAA CUGAUGAG X CGAA IGGCAGCA	933	TGCTGCCC A TTATACTC	1363
721	GCAUUGGG CUGAUGAG X CGAA IUUAUAUG	934	CATTATAC T CCCAATGC	1364
723	CGGCAUUG CUGAUGAG X CGAA IAGUAUAA	935	TTATACTC C CAATGCCG	1365
724	CCGGCAUU CUGAUGAG X CGAA IGAGUAUA	936	TATACTCC C AATGCCG	1366
725	ACCGGCAU CUGAUGAG X CGAA IGGAGUAU	937	ATACTCCC A ATGCCGGT	1367
730	GUGUCACC CUGAUGAG X CGAA ICAUUGGG	938	CCCAATGC C GGTGACAC	1368
737	UACUGUUG CUGAUGAG X CGAA IUCACCGG	939	CCGGTGAC A CAACAGTA	1369
739	AAUACUGU CUGAUGAG X CGAA IUGUCACC	940	GGTGACAC A ACAGTATT	1370
742	UGUAAUAC CUGAUGAG X CGAA IUUGUGUC	941	GACACAAC A GTATTACA	1371
750	CAUCAUAC CUGAUGAG X CGAA IUAAUACU	942	AGTATTAC A GTATGATG	1372
761	UUUACAGA CUGAUGAG X CGAA IUCAUCAU	943	ATGATGAC A TCTGTAAA	1373
764	UAUUUUAC CUGAUGAG X CGAA IAUGUCAU	944	ATGACATC T GTAAAATA	1374
776	UGUCCCAA CUGAUGAG X CGAA IUCUAUUU	945	AAATAGAC T TTGGAACA	1375
784	CUUAUAUG CUGAUGAG X CGAA IUUCCAAA	946	TTTGGAAC A CATATAAG	1376
786	CACUUAUA CUGAUGAG X CGAA IUGUCCAA	947	TGGAACAC A TATAAGTG	1377
809	AAAAGCAC CUGAUGAG X CGAA IUCAAUAA	948	TTATTGAC T GTGCTTTT	1378
814	ACAGUAAA CUGAUGAG X CGAA ICACAGUC	949	GACTGTGC T TTTACTGT	1379
820	AAAGUGAC CUGAUGAG X CGAA IUAAAAGC	950	GCTTTTAC T GTCACTTT	1380
824	AUUAUAAAG CUGAUGAG X CGAA IACAGUAA	951	TTACTGTC A CTTTTAAT	1381
826	GGAUUAUAA CUGAUGAG X CGAA IUGACAGU	952	ACTGTGTC T TTTAATCC	1382
834	CAUAUUUG CUGAUGAG X CGAA IAUUAAAA	953	TTTTAATC C CAAATATG	1383
835	UCAUAUUU CUGAUGAG X CGAA IGAUUAAA	954	TTTAATCC C AAATATGA	1384
836	AUCAUAUU CUGAUGAG X CGAA IGGAUUAA	955	TTAATCCC A AATATGAT	1385
859	UCUUUUAC CUGAUGAG X CGAA ICUUUUAA	956	TTAAAAGC T GTAAAAGA	1386
871	GUGUUAGU CUGAUGAG X CGAA ICAUCUUU	957	AAAGATGC T ACTAACAC	1387
874	CCAGUGUU CUGAUGAG X CGAA IUAGCAUC	958	GATGCTAC T AACACTGG	1388
878	UAUUCCAG CUGAUGAG X CGAA IUUAGUAG	959	CTACTAAC A CTGGAATA	1389
880	UUUAUUC CUGAUGAG X CGAA IUGUAGU	960	ACTAACAC T GGAATAAA	1390

Table 10

895	UCAAUCC CUGAUGAG X CGAA ICACACUU	961	AAGTGTGC T GGAATTGA	1391
912	CAUCACAC CUGAUGAG X CGAA IACGAACA	962	TGTTGTC T GTGTGATG	1392
931	UCUUGGAU CUGAUGAG X CGAA ICCUCACC	963	GGTGAGGC C ATCCAAGA	1393
932	UUCUUGGA CUGAUGAG X CGAA IGCCUCAC	964	GTGAGGCC A TCCAAGAA	1394
935	AACUUCUU CUGAUGAG X CGAA IAUGGCCU	965	AGGCCATC C AAGAAGTT	1395
936	UAACUUCU CUGAUGAG X CGAA IGAUGGCC	966	GGCCATCC A AGAAGTTA	1396
952	ACUUCAUA CUGAUGAG X CGAA IACUCCAU	967	ATGGAGTC C TATGAAGT	1397
953	AACUUCAU CUGAUGAG X CGAA IGACUCCA	968	TGGAGTCC T ATGAAGTT	1398
979	ACUUGAUA CUGAUGAG X CGAA IUCUCCCC	969	GGGAAGAC A TATCAAGT	1399
984	GUUUCACU CUGAUGAG X CGAA IAU AUGUC	970	GACATATC A AGTGAAAC	1400
993	UACGGAU CUGAUGAG X CGAA IUUUCACU	971	AGTGAAAC C AATCCGTA	1401
994	UUACGGAU CUGAUGAG X CGAA IGUUUCAC	972	GTGAAACC A ATCCGTAA	1402
998	UAGAUUAC CUGAUGAG X CGAA IAUUGGUU	973	AACCAATC C GTAATCTA	1403
1005	GUCCAUUU CUGAUGAG X CGAA IAUUACGG	974	CCGTAATC T AAATGGAC	1404
1014	CAAUUGAA CUGAUGAG X CGAA IUCCAUIU	975	AAATGGAC A TTCAATTG	1405
1018	UGCCCAAU CUGAUGAG X CGAA IAAUGUCC	976	GGACATTC A ATTGGGCA	1406
1026	UUCUAUAU CUGAUGAG X CGAA ICCCAAUU	977	AATTGGGC A ATATAGAA	1407
1038	UUCAGCA CUGAUGAG X CGAA IUAUUCUA	978	TAGAATAC A TGCTGGAA	1408
1042	GUUUUCC CUGAUGAG X CGAA ICAUGUAU	979	ATACATGC T GGAAAAAC	1409
1051	AUCGGCAC CUGAUGAG X CGAA IUUUUCC	980	GGAAAAAC A GTGCCGAT	1410
1056	UCACAAUC CUGAUGAG X CGAA ICACUGUU	981	AACAGTGC C GATTGTGA	1411
1078	AUUCUUGU CUGAUGAG X CGAA ICCUCCCC	982	GGGGAGGC A ACAAGAAT	1412
1081	UCCAUCU CUGAUGAG X CGAA IUUGCCUC	983	GAGGCAAC A AGAATGGA	1413
1108	GUUUCAAU CUGAUGAG X CGAA ICAUAUAC	984	GTATATGC A ATTGAAAC	1414
1117	CUACAAA CUGAUGAG X CGAA IUUUCAU	985	ATTGAAAC C TTTGGTAG	1415
1118	ACUACCAA CUGAUGAG X CGAA IGUUCAA	986	TTGAAACC T TTGGTAGT	1416
1129	CCUUUCC CUGAUGAG X CGAA IUACUACC	987	GGTAGTAC A GGAAAAGG	1417
1146	UAUCAUA CUGAUGAG X CGAA IAACAACA	988	TGTTGTTC A TGATGATA	1418
1165	AUGUAAUG CUGAUGAG X CGAA IAACAUUC	989	GAATGTTC A CATTACAT	1419
1167	UCAUGUAA CUGAUGAG X CGAA IUGAACAU	990	ATGTTTAC A TTACATGA	1420
1172	AUUUUUCA CUGAUGAG X CGAA IUAAUGUG	991	CACATTAC A TGAAAAAT	1421
1194	UUGGCACA CUGAUGAG X CGAA IUCCAACA	992	TGTTGGAC A TGTGCCAA	1422
1200	GCCUUAU CUGAUGAG X CGAA ICACAUGU	993	ACATGTGC C AATAAGGC	1423
1201	AGCCUUAU CUGAUGAG X CGAA IGCACAUG	994	CATGTGCC A ATAAGGCT	1424
1209	UUCUUGGA CUGAUGAG X CGAA ICCUUAU	995	AATAAGGC T TCCAAGAA	1425
1212	UUGUUCU CUGAUGAG X CGAA IAAGCCU	996	AAGGCTTC C AAGAACAA	1426
1213	UUUGUUCU CUGAUGAG X CGAA IGAAGCCU	997	AGGCTTCC A AGAACAAA	1427
1219	AAGUGUUU CUGAUGAG X CGAA IUUCUUGG	998	CCAAGAAC A AAACACTT	1428
1224	UUAACAAG CUGAUGAG X CGAA IUUUUGUU	999	AACAAAAC A CTTGTTAA	1429
1226	AUUUAACA CUGAUGAG X CGAA IUGUUUUG	1000	CAAAACAC T TGTTAAAT	1430
1238	UUCAUUGA CUGAUGAG X CGAA IACAUUUA	1001	TAAATGTC A TCAATGAA	1431
1241	GUUUUCAU CUGAUGAG X CGAA IAUGACAU	1002	ATGTCATC A ATGAAAAC	1432
1250	GGUCCAA CUGAUGAG X CGAA IUUUUCAU	1003	ATGAAAAC T TTGGAACC	1433
1258	AAGGCAAG CUGAUGAG X CGAA IUUCCAAA	1004	TTTGGAAC C CTTGCCTT	1434
1259	GAAGGCAA CUGAUGAG X CGAA IGUUCCAA	1005	TTGGAACC C TTGCCTTC	1435
1260	AGAAGGCA CUGAUGAG X CGAA IGGUCCAA	1006	TGGAACCC T TGCCTTCT	1436
1264	CGGCAGAA CUGAUGAG X CGAA ICAAGGGU	1007	ACCCTTGC C TTCTGCCG	1437

Table 10

1265	GCGGCAGA CUGAUGAG X CGAA IGCAAGGG	1008	CCCTTGCC T TCTGCCGC	1438
1268	UCUGCGGC CUGAUGAG X CGAA IAAGGCAA	1009	TTGCCTTC T GCCGCAGA	1439
1271	CCAUCUGC CUGAUGAG X CGAA ICAGAAGG	1010	CCTTCTGC C GCAGATGG	1440
1274	CAGCCAUC CUGAUGAG X CGAA ICGGCAGA	1011	TCTGCCGC A GATGGCTG	1441
1281	AGCGAUCC CUGAUGAG X CGAA ICCAUCUG	1012	CAGATGGC T GGATCGCT	1442
1289	UUCUCCCA CUGAUGAG X CGAA ICGAUCCA	1013	TGGATCGC T TGGGAGAA	1443
1307	AGCCAUCA CUGAUGAG X CGAA IUUUUAC	1014	GTAAATAC T TGATGGCT	1444
1315	UUCUUCAG CUGAUGAG X CGAA ICCAUCAA	1015	TTGATGGC T CTGAAGAA	1445
1317	GAUUCUUC CUGAUGAG X CGAA IAGCCAUC	1016	GATGGCTC T GAAGAATC	1446
1326	AGUCACAC CUGAUGAG X CGAA IAUUCUUC	1017	GAAGAATC T GTGTGACT	1447
1334	AAUGCCCA CUGAUGAG X CGAA IUCACACA	1018	TGTGTGAC T TGGGCATT	1448
1340	AUCUACAA CUGAUGAG X CGAA ICCCAAGU	1019	ACTTGGGC A TTGTAGAT	1449
1350	GUGGAUUA CUGAUGAG X CGAA IAUCUACA	1020	TGTAGATC C ATATCCAC	1450
1351	GGUGGAUA CUGAUGAG X CGAA IGAUCUAC	1021	GTAGATCC A TATCCACC	1451
1356	AUAAUGGU CUGAUGAG X CGAA IAUUGGA	1022	TCCATATC C ACCATTAT	1452
1357	CAUAAUGG CUGAUGAG X CGAA IGAUUGG	1023	CCATATCC A CCATTATG	1453
1359	CACAUAAU CUGAUGAG X CGAA IUUGGAUUA	1024	ATATCCAC C ATTATGTG	1454
1360	UCACAUAA CUGAUGAG X CGAA IGUGGAUA	1025	TATCCACC A TTATGTGA	1455
1370	UCCUUUAA CUGAUGAG X CGAA IUCACAU	1026	TATGTGAC A TTAAAGGA	1456
1381	GCUGUAUA CUGAUGAG X CGAA IAUCCUUU	1027	AAAGGATC A TATACAGC	1457
1387	AAUUGCGC CUGAUGAG X CGAA IUUAUGA	1028	TCATATAC A GCGCAATT	1458
1392	GUUCAAAU CUGAUGAG X CGAA ICGCUGUA	1029	TACAGCGC A ATTTGAAC	1459
1401	GGAUGGUA CUGAUGAG X CGAA IUUCAAU	1030	ATTTGAAC A TACCATCC	1460
1405	AACAGGAU CUGAUGAG X CGAA IUAGUUC	1031	GAACATAC C ATCCTGTT	1461
1406	CAACAGGA CUGAUGAG X CGAA IGUAUGUU	1032	AACATACC A TCCTGTTG	1462
1409	ACGCAACA CUGAUGAG X CGAA IAUUGUAU	1033	ATACCATC C TGTGCGT	1463
1410	GACGCAAC CUGAUGAG X CGAA IGAUGGUA	1034	TACCATCC T GTTGCGTC	1464
1419	UACAUGUU CUGAUGAG X CGAA IACGCAAC	1035	GTTGCGTC C AACATGTA	1465
1420	UUACAUGU CUGAUGAG X CGAA IGACGCAA	1036	TTGCGTCC A ACATGTAA	1466
1423	UCUUUACA CUGAUGAG X CGAA IUUGGACG	1037	CGTCCAAC A TGTAAGGA	1467
1439	UCCUCUGC CUGAUGAG X CGAA IACAACUU	1038	AAGTTGTC A GCAGAGGA	1468
1442	AUCUCCUC CUGAUGAG X CGAA ICUGACAA	1039	TTGTCAGC A GAGGAGAT	1469
1454	AGUUUAAU CUGAUGAG X CGAA IUCAUCUC	1040	GAGATGAC T ATTAAACT	1470
1462	UUGGACUA CUGAUGAG X CGAA IUUUAAUA	1041	TATTAAAC T TAGTCCAA	1471
1468	GUGGCUUU CUGAUGAG X CGAA IACUAAGU	1042	ACTTAGTC C AAAGCCAC	1472
1469	GGUGGCUU CUGAUGAG X CGAA IGACUAAG	1043	CTTAGTCC A AAGCCACC	1473
1474	GUUGAGGU CUGAUGAG X CGAA ICUUUGGA	1044	TCCAAAGC C ACCTCAAC	1474
1475	UGUUGAGG CUGAUGAG X CGAA IGCUUUGG	1045	CCAAAGCC A CCTCAACA	1475
1477	GGUGUUGA CUGAUGAG X CGAA IUUGCUUU	1046	AAAGCCAC C TCAACACC	1476
1478	AGGUGUUG CUGAUGAG X CGAA IGUGGCUU	1047	AAGCCACC T CAACACCT	1477
1480	AAAGGUGU CUGAUGAG X CGAA IAGGUGGC	1048	GCCACCTC A ACACCTTT	1478
1483	AAUAAAGG CUGAUGAG X CGAA IUUGAGGU	1049	ACCTCAAC A CCTTTATT	1479
1485	AAAAUAAA CUGAUGAG X CGAA IUGUUGAG	1050	CTCAACAC C TTTATTTT	1480
1486	GAAAAUAA CUGAUGAG X CGAA IGUGUUGA	1051	TCAACACC T TTATTTTC	1481
1495	CAAAGCUC CUGAUGAG X CGAA IAAAAUAA	1052	TTATTTTC T GAGCTTTG	1482
1500	UCCAACAA CUGAUGAG X CGAA ICUCAGAA	1053	TTCTGAGC T TTGTTGGA	1483
1513	UGGUAUCA CUGAUGAG X CGAA IUUUUCCA	1054	TGGAAAAC A TGATACCA	1484

Table 10

1520	UUAAUUCU CUGAUGAG X CGAA IUUAUCAUG	1055	CATGATAC C AGAATTAA	1485
1521	AUUAAUUC CUGAUGAG X CGAA IGUAUCAU	1056	ATGATACC A GAATTAAT	1486
1534	CAACAUGU CUGAUGAG X CGAA ICAAAUUA	1057	TAATTTGC C ACATGTTG	1487
1535	ACAACAUG CUGAUGAG X CGAA IGCAAAU	1058	AATTTGCC A CATGTTGT	1488
1537	AGACAACA CUGAUGAG X CGAA IUGGCAAA	1059	TTTGCCAC A TGTGTCT	1489
1545	GUUAAAAC CUGAUGAG X CGAA IACAACAU	1060	ATGTTGTC T GTTTTAAC	1490
1554	GGGUCCAC CUGAUGAG X CGAA IUUAAAAC	1061	GTTTTAAC A GTGGACCC	1491
1561	AUUACAUG CUGAUGAG X CGAA IUCCACUG	1062	CAGTGGAC C CATGTAAT	1492
1562	UAUUACAU CUGAUGAG X CGAA IGUCCACU	1063	AGTGGACC C ATGTAATA	1493
1563	GUUUACA CUGAUGAG X CGAA IGGUCCAC	1064	GTGGACCC A TGTAATAC	1494
1572	UGGAUAAA CUGAUGAG X CGAA IUUUACA	1065	TGTAATAC T TTTATCCA	1495
1579	UUAAACAU CUGAUGAG X CGAA IAUAAAAG	1066	CTTTTATC C ATGTTTAA	1496
1580	UUUAAAAC CUGAUGAG X CGAA IGAUAAAA	1067	TTTTATCC A TGTTTAAA	1497
1606	UUUGCCUU CUGAUGAG X CGAA IUCCAAAU	1068	ATTGGAC A AAGGCAAA	1498
1612	AGACGGUU CUGAUGAG X CGAA ICCUUUGU	1069	ACAAAGGC A AACCGTCT	1499
1616	CAUUAGAC CUGAUGAG X CGAA IUUUGCCU	1070	AGGCAAAC C GTCTAATG	1500
1620	AUUACAUI CUGAUGAG X CGAA IACGGUUU	1071	AAACCGTC T AATGTAAT	1501
1633	UUUUCGU CUGAUGAG X CGAA IUUAAUUA	1072	TAATTAAC C AACGAAAA	1502
1634	UUUUUCGU CUGAUGAG X CGAA IGUAAUUA	1073	AATTAACC A ACGAAAAA	1503
1645	GUCCGGAA CUGAUGAG X CGAA ICUUUUUC	1074	GAAAAAGC T TTCCGGAC	1504
1649	AAAAGUCC CUGAUGAG X CGAA IAAAGCUU	1075	AAGCTTTC C GGACTTTT	1505
1654	CAUUUAAA CUGAUGAG X CGAA IUCCGGAA	1076	TTCCGGAC T TTTAAATG	1506
1664	AAACAGUU CUGAUGAG X CGAA ICAUUUAA	1077	TTAAATGC T AACTGTTT	1507
1668	GGAAAAAC CUGAUGAG X CGAA IUUAGCAU	1078	ATGCTAAC T GTTTTTCC	1508
1676	CAGGAAGG CUGAUGAG X CGAA IAAAAACA	1079	TGTTTTTC C CCTTCCTG	1509
1677	ACAGGAAG CUGAUGAG X CGAA IGAAAAAC	1080	GTTTTTCC C CTTCCTGT	1510
1678	GACAGGAA CUGAUGAG X CGAA IGGAAAAA	1081	TTTTTCCC C TTCCTGTC	1511
1679	AGACAGGA CUGAUGAG X CGAA IGGGAAAA	1082	TTTTCCCC T TCCTGTCT	1512
1682	CCUAGACA CUGAUGAG X CGAA IAAGGGGA	1083	TCCCCTTC C TGTCTAGG	1513
1683	UCCUAGAC CUGAUGAG X CGAA IGAAGGGG	1084	CCCCTTCC T GTCTAGGA	1514
1687	AUUUUCCU CUGAUGAG X CGAA IACAGGAA	1085	TTCCTGTC T AGGAAAT	1515
1698	AGCUUUU CUGAUGAG X CGAA ICAUUUUC	1086	GAAATGTC T ATAAAGCT	1516
1706	CUAAUUUG CUGAUGAG X CGAA ICUUUUAU	1087	TATAAAGC T CAAATTAG	1517
1708	AACUAAUU CUGAUGAG X CGAA IAGCUUUA	1088	TAAAGCTC A AATTAGTT	1518
1726	AACGUUAU CUGAUGAG X CGAA IUCAUUC	1089	GGAATGAC T TATACGTT	1519
1748	AUCUCUUA CUGAUGAG X CGAA IUUAUCAA	1090	TTGAATAC C TAAGAGAT	1520
1749	UAUCUCUU CUGAUGAG X CGAA IGUAUUCA	1091	TGAATACC T AAGAGATA	1521
1759	AUCCAAAA CUGAUGAG X CGAA IUUAUCUCU	1092	AGAGATAC T TTTTGGAT	1522
1779	AAGAAUUA CUGAUGAG X CGAA ICAAUUAU	1093	TATATTGC C ATATTCTT	1523
1780	UAAGAAUA CUGAUGAG X CGAA IGCAAUUA	1094	ATATTGCC A TATTCTTA	1524
1786	UUCAAGUA CUGAUGAG X CGAA IAAUAUGG	1095	CCATATTC T TACTTGAA	1525
1790	AGCAUUCA CUGAUGAG X CGAA IUAGAAU	1096	ATTCTTAC T TGAATGCT	1526
1798	UCAUUCUA CUGAUGAG X CGAA ICAUUCUA	1097	TTGAATGC T TTGAATGA	1527
1808	CUGGAUGU CUGAUGAG X CGAA IUCAUUCA	1098	TGAATGAC T ACATCCAG	1528
1811	GAACUGGA CUGAUGAG X CGAA IUAGUCAU	1099	ATGACTAC A TCCAGTTC	1529
1814	GCAGAAU CUGAUGAG X CGAA IAUGUAGU	1100	ACTACATC C AGTCTGTC	1530
1815	UGCAGAAC CUGAUGAG X CGAA IGAUGUAG	1101	CTACATCC A GTTCTGCA	1531

Table 10

1820	AUAGGUGC CUGAUGAG X CGAA IAACUGGA	1102	TCCAGTTC T GCACCTAT	1532
1823	GGUAUAGG CUGAUGAG X CGAA ICAGAACU	1103	AGTTCTGC A CCTATACC	1533
1825	AGGGUUAU CUGAUGAG X CGAA IUGCAGAA	1104	TTCTGCAC C TATACCCT	1534
1826	GAGGGUUAU CUGAUGAG X CGAA IGUCAGAA	1105	TCTGCACC T ATACCCTC	1535
1831	CACCAGAG CUGAUGAG X CGAA IUUAUAGGU	1106	ACCTATAC C CTCTGGTG	1536
1832	ACACCAGA CUGAUGAG X CGAA IGUAUAGG	1107	CCTATACC C TCTGGTGT	1537
1833	AACACCAG CUGAUGAG X CGAA IGGUAUAG	1108	CTATACCC T CTGGTGTT	1538
1835	GCAACACC CUGAUGAG X CGAA IAGGGUUAU	1109	ATACCCTC T GGTGTTGC	1539
1844	GGUUAUAA CUGAUGAG X CGAA ICAACACC	1110	GGTGTGTC T TTTTAACC	1540
1852	UCCAGGAA CUGAUGAG X CGAA IUUAAAAA	1111	TTTTTAAC C TTCCTGGA	1541
1853	UUCAGGAA CUGAUGAG X CGAA IGUAAAAA	1112	TTTTAACC T TCCTGGAA	1542
1856	GGAUUCCA CUGAUGAG X CGAA IAAGGUUA	1113	TAACCTTC C TGGAATCC	1543
1857	UGGAUCC CUGAUGAG X CGAA IGAAGGUU	1114	AACCTTCC T GGAATCCA	1544
1864	UAGAAAAU CUGAUGAG X CGAA IAUUCCAG	1115	CTGGAATC C ATTTCTA	1545
1865	UUAGAAAA CUGAUGAG X CGAA IGAUCCA	1116	TGGAATCC A TTTCTAA	1546
1871	UAUUUUUU CUGAUGAG X CGAA IAAAUGG	1117	CCATTTTC T AAAAAATA	1547
1885	GAAGAAUG CUGAUGAG X CGAA IUCUUUAU	1118	ATAAGAC A CATCTTC	1548
1887	GAGAAGAA CUGAUGAG X CGAA IUGUCUUU	1119	AAAGACAC A TTCTCTC	1549
1891	UGCUGAGA CUGAUGAG X CGAA IAAUGUGU	1120	ACACATTC T TCTCAGCA	1550
1894	UGGUGCUG CUGAUGAG X CGAA IAAGAAUG	1121	CATTCTTC T CAGACCA	1551
1896	UGUGGUGC CUGAUGAG X CGAA IAGAAGAA	1122	TTCTTCTC A GCACCACA	1552
1899	UUGUGUGG CUGAUGAG X CGAA ICUGAGAA	1123	TTCTCAGC A CCACACAA	1553
1901	UGUUGUGU CUGAUGAG X CGAA IUGCUGAG	1124	CTCAGCAC C ACACAACA	1554
1902	GUGUUGUG CUGAUGAG X CGAA IGUCUGA	1125	TCAGCACC A CACAACAC	1555
1904	AGGUGUUG CUGAUGAG X CGAA IUGGUGCU	1126	AGCACCAC A CAACACCT	1556
1906	AUAGGUGU CUGAUGAG X CGAA IUGUGGUG	1127	CACCACAC A ACACCTAT	1557
1909	GGAAUAGG CUGAUGAG X CGAA IUUGUGUG	1128	CACACAAC A CCTATTCC	1558
1911	UUGGAAUA CUGAUGAG X CGAA IUGUUGUG	1129	CACAACAC C TATTCCAA	1559
1912	UUUGGAAU CUGAUGAG X CGAA IGUGUUGU	1130	ACAACACC T ATTCCAAA	1560
1917	UCGAUUUU CUGAUGAG X CGAA IAAUAGGU	1131	ACCTATTC C AAAATCGA	1561
1918	GUCGAUUU CUGAUGAG X CGAA IGAUAGG	1132	CCTATTCC A AAATCGAC	1562
1927	AAAUUUGU CUGAUGAG X CGAA IUCGAUUU	1133	AAATCGAC C ACATATTT	1563
1928	CAAAUAUG CUGAUGAG X CGAA IGUCGAUU	1134	AATCGACC A CATATTTG	1564
1930	UCCAAAUA CUGAUGAG X CGAA IUGGUCGA	1135	TCGACCAC A TATTTGGA	1565
1947	CUGAGGAG CUGAUGAG X CGAA ICUUUAU	1136	AGTAAAGC T CTCCTCAG	1566
1949	UGCUGAGG CUGAUGAG X CGAA IAGCUUUA	1137	TAAAGCTC T CCTCAGCA	1567
1951	UUUGCUGA CUGAUGAG X CGAA IAGAGCUU	1138	AAGCTCTC C TCAGCAAA	1568
1952	AUUUGCUG CUGAUGAG X CGAA IGAGAGCU	1139	AGCTCTCC T CAGCAAAT	1569
1954	ACAUUUGC CUGAUGAG X CGAA IAGGAGAG	1140	CTCTCCTC A GCAAATGT	1570
1957	UUUACAUU CUGAUGAG X CGAA ICUGAGGA	1141	TCCTCAGC A AATGTAAA	1571
1971	AUAAUUUC CUGAUGAG X CGAA IUUCUUUU	1142	AAAAGAAC A GAAATTAT	1572
1983	AGACAGUU CUGAUGAG X CGAA IUUAUAAU	1143	ATTATAAC A AACTGTCT	1573
1987	UGAGAGAC CUGAUGAG X CGAA IUUUGUUA	1144	TAACAAAC T GTCTCTCA	1574
1991	GGUCUGAG CUGAUGAG X CGAA IACAGUUU	1145	AAACTGTC T CTCAGACC	1575
1993	GUGGUCUG CUGAUGAG X CGAA IAGACAGU	1146	ACTGTCTC T CAGACCAC	1576
1995	CUGUGGUC CUGAUGAG X CGAA IAGAGACA	1147	TGTCTCTC A GACCACAG	1577
1999	UAUACUGU CUGAUGAG X CGAA IUCUGAGA	1148	TCTCAGAC C ACAGTATA	1578

Table 10

2000	UUUACUG CUGAUGAG X CGAA IGUCUGAG	1149	CTCAGACC A CAGTATAA	1579
2002	GGUUUAC CUGAUGAG X CGAA IUGGUCUG	1150	CAGACCAC A GTATAACC	1580
2010	UCUAGUUU CUGAUGAG X CGAA IUUUAUCU	1151	AGTATAAC C AAAC TAGA	1581
2011	UUCUAGUU CUGAUGAG X CGAA IGUUAUAC	1152	GTATAACC A AACTAGAA	1582
2015	UGAGUUCU CUGAUGAG X CGAA IUUUGGUU	1153	AACCAAAC T AGAACTCA	1583
2021	UAAUCCUG CUGAUGAG X CGAA IUUCUAGU	1154	ACTAGAAC T CAGGATTA	1584
2023	CUUAAUCC CUGAUGAG X CGAA IAGUUCUA	1155	TAGAACTC A GGATTAAG	1585
2036	UUUGAGUG CUGAUGAG X CGAA IUUUCUUA	1156	TAAGAAAC T CACTCAA	1586
2038	GUUUUGAG CUGAUGAG X CGAA IAGUUUCU	1157	AGAACTC A CTCAAAC	1587
2040	UGGUUUUG CUGAUGAG X CGAA IUGAGUUU	1158	AAACTCAC T CAAAACCA	1588
2042	UGUGUUUU CUGAUGAG X CGAA IAGUGAGU	1159	ACTCACTC A AAACCACA	1589
2047	AGUUGUGU CUGAUGAG X CGAA IUUUUGAG	1160	CTCAAAC C ACACAACT	1590
2048	UAGUUGUG CUGAUGAG X CGAA IGUUUUGA	1161	TCAAAC C CACAATA	1591
2050	UGUAGUUG CUGAUGAG X CGAA IUGGUUUU	1162	AAAACCAC A CAACTACA	1592
2052	CAUGUAGU CUGAUGAG X CGAA IUGUGGUU	1163	AACCACAC A ACTACATG	1593
2055	UUCCAUGU CUGAUGAG X CGAA IUUGUGUG	1164	CACACAAC T ACATGGAA	1594
2058	AGUUUCCA CUGAUGAG X CGAA IUAGUUGU	1165	ACAAC TAC A TGGAACT	1595
2066	GGUUGUUC CUGAUGAG X CGAA IUUCCAU	1166	ATGGAAAC T GAACAACC	1596
2071	GAGCAGGU CUGAUGAG X CGAA IUUCAGUU	1167	AACTGAAC A ACCTGCTC	1597
2074	CAGGAGCA CUGAUGAG X CGAA IUUGUUCA	1168	TGAACAAC C TGCTCCTG	1598
2075	UCAGGAGC CUGAUGAG X CGAA IGUUGUUC	1169	GAACAACC T GCTCCTGA	1599
2078	CAUUCAGG CUGAUGAG X CGAA ICAGGUUG	1170	CAACCTGC T CCTGAATG	1600
2080	GUCAUUC CUGAUGAG X CGAA IAGCAGGU	1171	ACCTGCTC C TGAATGAC	1601
2081	AGUCAUUC CUGAUGAG X CGAA IAGCAGG	1172	CCTGCTCC T GAATGACT	1602
2089	UAUCCAGU CUGAUGAG X CGAA IUCAUUC	1173	TGAATGAC T ACTGGATA	1603
2092	AUGUAUCC CUGAUGAG X CGAA IUAGUCAU	1174	ATGACTAC T GGATACAT	1604
2099	UUUUGUUA CUGAUGAG X CGAA IUUAUCCAG	1175	CTGGATAC A TAACAAA	1605
2104	CUUCAUUU CUGAUGAG X CGAA IUUAUGUA	1176	TACATAAC A AAATGAAG	1606
2115	UUUAUUUC CUGAUGAG X CGAA ICCUUCAU	1177	ATGAAGGC A GAAATAA	1607
2131	GGUUUUA CUGAUGAG X CGAA IAAAUUCU	1178	AGATGTTC T TTAACACC	1608
2139	UUCUCAUU CUGAUGAG X CGAA IUUUUAAA	1179	TTTAAAC C AATGAGAA	1609
2140	GUUCUCAU CUGAUGAG X CGAA IGUUUUA	1180	TTAAAC C AATGAGAA	1610
2149	UGUGUCUU CUGAUGAG X CGAA IUUCUCAU	1181	ATGAGAAC A AAGACACA	1611
2155	GUAGUUG CUGAUGAG X CGAA IUCUUGU	1182	ACAAAGAC A CAACATAC	1612
2157	UGGUAGU CUGAUGAG X CGAA IUGUCUUU	1183	AAAGACAC A ACATACCA	1613
2160	UUCUGGUA CUGAUGAG X CGAA IUUGUGUC	1184	GACACAAC A TACCAGAA	1614
2164	GAGAUUCU CUGAUGAG X CGAA IUUGUUG	1185	CAACATAC C AGAATCTC	1615
2165	AGAGAUUC CUGAUGAG X CGAA IGUAUGUU	1186	AACATACC A GAATCTCT	1616
2171	UGUCCAG CUGAUGAG X CGAA IAUUCUGG	1187	CCAGAATC T CTGGGACA	1617
2173	UGUGUCC CUGAUGAG X CGAA IAGAUUCU	1188	AGAATCTC T GGGACACA	1618
2179	UUUGAAUG CUGAUGAG X CGAA IUCCAGAA	1189	TCTGGGAC A CATTCAAA	1619
2181	GCUUGAA CUGAUGAG X CGAA IUGUCCA	1190	TGGGACAC A TTCAAAGC	1620
2185	CACUGCUU CUGAUGAG X CGAA IAAUGUGU	1191	ACACATTC A AAGCAGTG	1621
2190	CUACACAC CUGAUGAG X CGAA ICUUGAA	1192	TTCAAAGC A GTGTGTAG	1622
2214	GCAUUUAG CUGAUGAG X CGAA ICUUAAA	1193	TTTATAGC A CTAAATGC	1623
2216	GGGAUUU CUGAUGAG X CGAA IUGCUAUA	1194	TATAGCAC T AAATGCCC	1624
2223	CUCUUGUG CUGAUGAG X CGAA ICAUUUAG	1195	CTAAATGC C CACAAGAG	1625

Table 10

2224	UCUCUUGU CUGAUGAG X CGAA IGCAUUA	1196	TAAATGCC C ACAAGAGA	1626
2225	UUCUCUUG CUGAUGAG X CGAA IGGCAUUU	1197	AAATGCCC A CAAGAGAA	1627
2227	CUUUCUCU CUGAUGAG X CGAA IUGGGCAU	1198	ATGCCAC A AGAGAAAG	1628
2237	AUAUUUCC CUGAUGAG X CGAA ICUUUCUC	1199	GAGAAAGC A GGAATAT	1629
2247	UCAUUUUU CUGAUGAG X CGAA IAUUUUUC	1200	GAAATATC T AAAATTGA	1630
2257	UGUUAGGG CUGAUGAG X CGAA IUCAAUUU	1201	AAATTGAC A CCCTAACA	1631
2259	GAUGUAG CUGAUGAG X CGAA IUGUCAAU	1202	ATTGACAC C CTAACATC	1632
2260	UGAUGUUA CUGAUGAG X CGAA IGUGUCAA	1203	TTGACACC C TAACATCA	1633
2261	GUGAUGUU CUGAUGAG X CGAA IGGUGUCA	1204	TGACACCC T AACATCAC	1634
2265	AAUUGUGA CUGAUGAG X CGAA IUUAGGGU	1205	ACCCTAAC A TCACAATT	1635
2268	UUUAUUG CUGAUGAG X CGAA IAUGUUAG	1206	CTAACATC A CAATTAAA	1636
2270	CUUUUAU CUGAUGAG X CGAA IUGAUGUU	1207	AACATCAC A ATTAAGA	1637
2282	GCUUCUCU CUGAUGAG X CGAA IUUCUUUU	1208	AAAAGAAC T AGAGAAGC	1638
2291	UUUGCUCU CUGAUGAG X CGAA ICUUCUCU	1209	AGAGAAGC A AGAGCAA	1639
2297	AAUGUGUU CUGAUGAG X CGAA ICUCUUGC	1210	GCAAGAGC A AACACATT	1640
2301	UUUCAUUG CUGAUGAG X CGAA IUUUGCUC	1211	GAGCAAAC A CATTGAAA	1641
2303	CUUUUCA CUGAUGAG X CGAA IUGUUUGC	1212	GCAAACAC A TTGAAAAG	1642
2313	CUUCUCUU CUGAUGAG X CGAA ICUUUUCA	1213	TGAAAAGC T AAGAGAAG	1643
2324	UUAUUUCU CUGAUGAG X CGAA ICCUUCUC	1214	GAGAAGGC A AGAAATA	1644
2334	CUGAUCUU CUGAUGAG X CGAA IUUAUUUC	1215	GAAATAAC T AAGATCAG	1645
2341	UUCUGCUC CUGAUGAG X CGAA IAUCUUAG	1216	CTAAGATC A GAGCAGAA	1646
2346	UUCAGUUC CUGAUGAG X CGAA ICUCUGAU	1217	ATCAGAGC A GAACTGAA	1647
2351	UUUCCUUC CUGAUGAG X CGAA IUUCUGCU	1218	AGCAGAAC T GAAGGAAA	1648
2367	GUUUUUUG CUGAUGAG X CGAA IUCUCUUA	1219	ATAGAGAC A CAAAAAAC	1649
2369	GAGUUUUU CUGAUGAG X CGAA IUGUCUCU	1220	AGAGACAC A AAAAATC	1650
2376	UUUUGAAG CUGAUGAG X CGAA IUUUUUUG	1221	CAAAAAAC T CTTCAAAA	1651
2378	UUUUUGA CUGAUGAG X CGAA IAGUUUUU	1222	AAAAATC T TCAAAAAA	1652
2381	UGAUUUUU CUGAUGAG X CGAA IAAGAGUU	1223	AACTCTC A AAAAATCA	1653
2389	GGAUUCAU CUGAUGAG X CGAA IAUUUUUU	1224	AAAAATC A ATGAATCC	1654
2397	CAGCUCCU CUGAUGAG X CGAA IAUUCAUU	1225	AATGAATC C AGGAGCTG	1655
2398	CCAGCUCC CUGAUGAG X CGAA IGAUUCAU	1226	ATGAATCC A GGAGCTGG	1656
2404	AAAAAAC CUGAUGAG X CGAA ICUCUGG	1227	CCAGGAGC T GGTTTTTT	1657
2422	AAUUUGU CUGAUGAG X CGAA IAUUGUUU	1228	AAACGATC A ACAAAT	1658
2425	AUCAUUU CUGAUGAG X CGAA IUUGAUCG	1229	CGATCAAC A AAATTGAT	1659
2438	CUUGCUG CUGAUGAG X CGAA IUCUAUCA	1230	TGATAGAC A CTAGCAAG	1660
2440	GUCUUGCU CUGAUGAG X CGAA IUGUCUUA	1231	ATAGACAC T AGCAAGAC	1661
2444	AUUAGUCU CUGAUGAG X CGAA ICUAGUGU	1232	ACACTAGC A AGACTAAT	1662
2449	UCUUUAUU CUGAUGAG X CGAA IUCUUGCU	1233	AGCAAGAC T AATAAGA	1663
2476	CUUCUAUU CUGAUGAG X CGAA IAUUCUUC	1234	GAAGAATC A AATAGAAG	1664
2486	UUUUUUAU CUGAUGAG X CGAA ICUUCUUA	1235	ATAGAAGC A ATAAAAA	1665
2511	AUUGGUGG CUGAUGAG X CGAA IAUUCC	1236	GGGATATC A CCACCAAT	1666
2513	GGAUUGGU CUGAUGAG X CGAA IUGAUUUC	1237	GATATCAC C ACCAATCC	1667
2514	GGGAUUGG CUGAUGAG X CGAA IGUGAUUU	1238	ATATCACC A CCAATCCC	1668
2516	GUGGGAUU CUGAUGAG X CGAA IUGUGUAU	1239	ATCACCAC C AATCCCAC	1669
2517	UGUGGGAU CUGAUGAG X CGAA IGUGGUGA	1240	TCACCACC A ATCCCACA	1670
2521	UUUCUGUG CUGAUGAG X CGAA IAUUGGUG	1241	CACCAATC C CACAGAAA	1671
2522	AUUUCUGU CUGAUGAG X CGAA IGAUUGGU	1242	ACCAATCC C ACAGAAAT	1672

Table 10

2523	UAUUUCUG CUGAUGAG X CGAA IGGAUUGG	1243	CCAATCCC A CAGAAATA	1673
2525	UUUAUUUC CUGAUGAG X CGAA IUGGGAUU	1244	AATCCCAC A GAAATAAA	1674
2535	CUGAUGGU CUGAUGAG X CGAA IUUUUAUU	1245	AAATAAAC C ACCATCAG	1675
2536	UCUGAUGG CUGAUGAG X CGAA IGUUUAUU	1246	AATAAACC A CCATCAGA	1676
2538	UCUCUGAU CUGAUGAG X CGAA IUGGUUUA	1247	TAAACCAC C ATCAGAGA	1677
2539	UUCUCUGA CUGAUGAG X CGAA IGUGGUUU	1248	AAACCACC A TCAGAGAA	1678
2542	GUAUUCUC CUGAUGAG X CGAA IAUGGUGG	1249	CCACCATC A GAGAATAC	1679
2551	GUGUUUGU CUGAUGAG X CGAA IUUUUCUC	1250	GAGAATAC T ACAAACAC	1680
2554	GAGGUGUU CUGAUGAG X CGAA IUAGUAUU	1251	AATACTAC A AACACCTC	1681
2558	CGUAGAGG CUGAUGAG X CGAA IUUUGUAG	1252	CTACAAAC A CCTCTACG	1682
2560	UGCGUAGA CUGAUGAG X CGAA IUGUUUGU	1253	ACAAACAC C TCTACGCA	1683
2561	UUGCGUAG CUGAUGAG X CGAA IGUGUUUG	1254	CAAACACC T CTACGCAA	1684

Input Sequence = HSU29607. Cut Site = CH/			
Stem Length = 8. Core Sequence = CUGAUGAG X CGAA (X = GCCGUUAGGC or other stem II)			
Seq1 = HSU29607 (Human methionine aminopeptidase mRNA, complete cds., 2569 bp)			

Table 11

Table 11: Human Methionine Aminopeptidase type 2 (MetAP-2) G-cleaver Ribozyme and Target Sequence

Nt. position	Substrate Sequence	Seq. ID Nos.	Ribozyme Sequence	Seq. ID Nos.
64	GGAGCCACCU G AAUGG	1685	CCAUU UGAUGGCAUGCACUAUGCGG AGGUGGCCUCC	1834
71	CCUGAAUGGC G ACCUG	1686	CAGGU UGAUGGCAUGCACUAUGCGG GCCAUUCAGG	1835
86	GGAUCCAGAC G ACAGG	1687	CCUGU UGAUGGCAUGCACUAUGCGG GUCUGGAUCC	1836
104	AGAAGGAGCU G CCUCU	1688	AGAGG UGAUGGCAUGCACUAUGCGG AGCUCCUUCU	1837
116	CUCUACGGCU G AGGAA	1689	UUCUU UGAUGGCAUGCACUAUGCGG AGCCGUAGAG	1838
141	AAAAAAGAC G AAAGA	1690	UCUUU UGAUGGCAUGCACUAUGCGG GUCUUUUUUU	1839
170	AGGGCCUUCU G CAGCA	1691	UGCUG UGAUGGCAUGCACUAUGCGG AGAAGGCCCU	1840
191	ACAGGAACCU G AUAAA	1692	UUUUU UGAUGGCAUGCACUAUGCGG AGGUUCCUGU	1841
218	CUCAGUGGAU G AAGUA	1693	UACUU UGAUGGCAUGCACUAUGCGG AUCCACUAGAG	1842
269	AGAAAGAGAU G AAGAU	1694	AUCUU UGAUGGCAUGCACUAUGCGG AUCUCUUUCU	1843
275	AGAUGAAGAU G AUGAA	1695	UUCAU UGAUGGCAUGCACUAUGCGG AUCUUCAUCU	1844
278	UGAAGAUGAU G AAGAU	1696	AUCUU UGAUGGCAUGCACUAUGCGG AUCAUCUCCA	1845
293	UGGAGAUGGC G AUGGA	1697	UCCAU UGAUGGCAUGCACUAUGCGG GCCAUCUCCA	1846
384	GUUCCAUAU G UGACC	1698	GGUCA UGAUGGCAUGCACUAUGCGG AUAUTUGGAAC	1847
386	UCCAUAUGU G ACCUG	1699	CAGGU UGAUGGCAUGCACUAUGCGG ACAUAUUGGA	1848
391	UAUGUGACCU G UAUCC	1700	GGAUA UGAUGGCAUGCACUAUGCGG AGGUCACAUAA	1849
404	UCCUAUUGGU G UAUUU	1701	AAAUU UGAUGGCAUGCACUAUGCGG ACCAUUAGGA	1850
426	GGACAAGAAU G CGAAU	1702	AUUUG UGAUGGCAUGCACUAUGCGG AUUCUUGUCC	1851
428	ACAAGAAGUC G AAUAC	1703	GUUUU UGAUGGCAUGCACUAUGCGG GCAUUCUUGU	1852
453	CAAGAUGGUC G AACAG	1704	CUUUU UGAUGGCAUGCACUAUGCGG GCCCAUCUUG	1853
461	GCGAACACGU G CUUGG	1705	CCAAG UGAUGGCAUGCACUAUGCGG AGCUUUGGC	1854
479	AACUACAAGU G AAGAA	1706	UUUUU UGAUGGCAUGCACUAUGCGG ACUUGUAGUU	1855
509	UCAGGCAAGU G AAGAG	1707	CUUUU UGAUGGCAUGCACUAUGCGG ACUUGCCUGA	1856
524	GAUUUGGAUU G AUUUU	1708	AAAAU UGAUGGCAUGCACUAUGCGG AUUCCAAAUC	1857
531	AAUGAUUUUC G AGAAG	1709	CUUUU UGAUGGCAUGCACUAUGCGG GAAAAUCAUU	1858
539	UCGAGAAGCU G CAGAA	1710	UUUUU UGAUGGCAUGCACUAUGCGG AGCUUCUCGA	1859
552	GAAGCACAUUC G ACAAG	1711	CUUUU UGAUGGCAUGCACUAUGCGG GAUGUGCUUC	1860
574	AAUACGUAAU G AGCUG	1712	CAGCU UGAUGGCAUGCACUAUGCGG AUUACGUAAU	1861

Table 11

595	AGCCUGGGAU G ACAAU	1713	AUUGU UGAUGGCAUGCACAUAUGCGG AUCCAGGCU	1862
601	GGAUGACAAU G AUAGA	1714	UCUAU UGAUGGCAUGCACAUAUGCGG AUUGUCAUC	1863
612	AUAGAAAUCU G UGAAA	1715	UUUCA UGAUGGCAUGCACAUAUGCGG AGAUUUUAU	1864
614	AGAAAUCUGU G AAAAG	1716	CUUUU UGAUGGCAUGCACAUAUGCGG ACAGAUUUU	1865
630	UUGGAAGACU G UUCAC	1717	GUGAA UGAUGGCAUGCACAUAUGCGG AGUCUUCCAA	1866
636	GACUGUUCAC G CAAGU	1718	ACUUG UGAUGGCAUGCACAUAUGCGG GUGAACAGUC	1867
665	UGGAUUAAAU G CAGGC	1719	GCCUG UGAUGGCAUGCACAUAUGCGG AUUUAAUCCA	1868
690	CCUACUGGAU G UUCUC	1720	GAGAA UGAUGGCAUGCACAUAUGCGG AUCCAGUAGG	1869
705	CUCAAUAUU G UGCUG	1721	CAGCA UGAUGGCAUGCACAUAUGCGG AAUUAUUGAG	1870
707	CAUAUAUUGU G CUGCC	1722	GGCAG UGAUGGCAUGCACAUAUGCGG ACAAUUAUUG	1871
710	UAAUUGUGU G CCCAU	1723	AUGGG UGAUGGCAUGCACAUAUGCGG AGCACAUAUA	1872
728	UACUCCCAU G CCGGU	1724	ACCGG UGAUGGCAUGCACAUAUGCGG AUUGGGAGUA	1873
734	CAUUGCCGGU G ACACA	1725	UGUGU UGAUGGCAUGCACAUAUGCGG ACCGGCAUUG	1874
755	AUUACAGUAU G AUGAC	1726	GUCAU UGAUGGCAUGCACAUAUGCGG AUACUGUAUU	1875
758	ACAGUAUGAU G ACAUC	1727	GAUGU UGAUGGCAUGCACAUAUGCGG AUCAUAUCUGU	1876
765	GAUGACAUCU G UAAAA	1728	UUUUA UGAUGGCAUGCACAUAUGCGG AGAUGUCAUC	1877
806	UAGGAUUAU G ACUGU	1729	ACAGU UGAUGGCAUGCACAUAUGCGG AAUAAUCCUA	1878
810	AUAUAUGACU G UGCUU	1730	AAGCA UGAUGGCAUGCACAUAUGCGG AGUCAUAUU	1879
812	UAUUGACUGU G CUUUU	1731	AAAAG UGAUGGCAUGCACAUAUGCGG ACAGUCAUA	1880
821	UGCUIUUACU G UCACU	1732	AGUGA UGAUGGCAUGCACAUAUGCGG AGUAAAAGCA	1881
842	UCCCAAAUAU G AUACG	1733	CGUAU UGAUGGCAUGCACAUAUGCGG AUUUUUGGGA	1882
860	AUUAAAAGCU G UAAAA	1734	UUUUA UGAUGGCAUGCACAUAUGCGG AGCUUUUAUU	1883
869	UGUAAAAGAU G CUACU	1735	AGUAG UGAUGGCAUGCACAUAUGCGG AUCUUUUACA	1884
891	GGAAUAAGU G UGCUG	1736	CAGCA UGAUGGCAUGCACAUAUGCGG ACUUUAUUC	1885
893	AAUAAAAGUGU G CUGGA	1737	UCCAG UGAUGGCAUGCACAUAUGCGG ACACUUUAUU	1886
902	UGCUGGAUUU G AUGUU	1738	AACAU UGAUGGCAUGCACAUAUGCGG AAUUCACGCA	1887
905	UGGAUUGAU G UUCGU	1739	ACGAA UGAUGGCAUGCACAUAUGCGG AUCAAUUCUA	1888
913	AUGUUGUCU G UGUGA	1740	UCACA UGAUGGCAUGCACAUAUGCGG AGACGAACAU	1889
915	GUUCGUCUGU G UGAUG	1741	CAUCA UGAUGGCAUGCACAUAUGCGG ACAGACGAAC	1890
917	UCGUCUGUGU G AUGUU	1742	AACAU UGAUGGCAUGCACAUAUGCGG ACACAGACGA	1891
920	UCUGUGUGAU G UUGGU	1743	ACCAA UGAUGGCAUGCACAUAUGCGG AUCAACACAGA	1892

Table 11

926	UGAUGUUGGU G AGGCC	1744	GGCCU UGAUGGCAUGCACUAUGCGCG ACCAACAUCA	1893
956	GGAGUCCUUAU G AAGUU	1745	AACUU UGAUGGCAUGCACUAUGCGCG AUAGGACUCC	1894
962	CUAUGAAGUU G AAUA	1746	UAUUU UGAUGGCAUGCACUAUGCGCG AACUUAUAG	1895
988	CAUAUCAAGU G AAACC	1747	GGUUU UGAUGGCAUGCACUAUGCGCG ACUUGAUUG	1896
1040	UAGAAUACAU G CUGGA	1748	UCCAG UGAUGGCAUGCACUAUGCGCG AUGUAUUCUA	1897
1054	GAAACACAGU G CCGAU	1749	AUCGG UGAUGGCAUGCACUAUGCGCG ACUGUUUUUC	1898
1057	AAACAGUGCC G AUUGU	1750	ACAAU UGAUGGCAUGCACUAUGCGCG GGCACUGUUU	1899
1061	AGUGCCGAUU G UGAAA	1751	UUUCA UGAUGGCAUGCACUAUGCGCG AAUCGGCACU	1900
1063	UGCCGAUUUG G AAAGG	1752	CCUUU UGAUGGCAUGCACUAUGCGCG ACAAU'CGCA	1901
1106	AGAAGUAUAU G CAAUU	1753	AAUUG UGAUGGCAUGCACUAUGCGCG AUAUACUUCU	1902
1112	AUAUGCAAUU G AAACC	1754	GGUUU UGAUGGCAUGCACUAUGCGCG AAUUGCAUAU	1903
1139	AGGAAAAGGU G UUGUU	1755	AACAA UGAUGGCAUGCACUAUGCGCG ACCUUUUCCU	1904
1142	AAAAGGUGUU G UUCAU	1756	AUGAA UGAUGGCAUGCACUAUGCGCG AACACCUUUU	1905
1148	UGUUGUUCAU G AUGAU	1757	AUCAU UGAUGGCAUGCACUAUGCGCG AUGAACAAACA	1906
1151	UGUUAUGAU G AUAUG	1758	CAUAU UGAUGGCAUGCACUAUGCGCG AUCAUGAACA	1907
1161	GAUAUGGAU G UUCAC	1759	GUGAA UGAUGGCAUGCACUAUGCGCG AUUCCAUAUC	1908
1174	CACAUUACAU G AAAAA	1760	UUUUU UGAUGGCAUGCACUAUGCGCG AUGUAAUGUG	1909
1184	GAAAAUUUU G AUGUU	1761	AACAU UGAUGGCAUGCACUAUGCGCG AAAAAUUUUU	1910
1187	AAAUUUUGAU G UUGGA	1762	UCCAA UGAUGGCAUGCACUAUGCGCG AUCAAAAUUU	1911
1196	UGUUGGACAU G UGCCA	1763	UGGCA UGAUGGCAUGCACUAUGCGCG AUGUCCAACA	1912
1198	UUGGACAUGU G CCAAU	1764	AUUGG UGAUGGCAUGCACUAUGCGCG ACAUGUCCAA	1913
1228	CAAAACACUU G UUAUA	1765	UUUAA UGAUGGCAUGCACUAUGCGCG AAGUGUUUUG	1914
1235	CUUGUUAAAU G UCAUC	1766	GAUGA UGAUGGCAUGCACUAUGCGCG AUUUAACAAG	1915
1244	UGUCAUCAAU G AAAC	1767	GUUUU UGAUGGCAUGCACUAUGCGCG AUUGAUGACA	1916
1262	UGGAACCCUU G CCUUC	1768	GAAGG UGAUGGCAUGCACUAUGCGCG AAGGGUUCCA	1917
1269	CUUGCCUUUC G CCGCA	1769	UGCGG UGAUGGCAUGCACUAUGCGCG AGAAGGCAAG	1918
1272	GCCUUCUGCC G CAGAU	1770	AUCUG UGAUGGCAUGCACUAUGCGCG GGCAGAAAGC	1919
1287	UGGCUGGAUC G CUUGG	1771	CCAAG UGAUGGCAUGCACUAUGCGCG GAUCCAGCCA	1920
1309	GUAAAUAUUU G AUGGC	1772	GCCAU UGAUGGCAUGCACUAUGCGCG AAGUAUUUAC	1921
1318	UGAUGGCUCU G AAGAA	1773	UUUUU UGAUGGCAUGCACUAUGCGCG AGAGCCAACA	1922
1327	UGAAGAAUUCU G UUGUA	1774	UCACA UGAUGGCAUGCACUAUGCGCG AGAUUCUUA	1923

Table 11

1329	AAGAAUCUGU G UGACU	1775	AGUCA UGAUGGCAUGCACAUAUGCGG ACAGAUUCUU	1924
1331	GAUCUGUGU G ACUUG	1776	CAAGU UGAUGGCAUGCACAUAUGCGG ACACAGAUUC	1925
1343	CUUGGGCAUU G UAGAU	1777	AUCUA UGAUGGCAUGCACAUAUGCGG AAUGCCCAAG	1926
1365	CCACCAUUAU G UGACA	1778	UGUCA UGAUGGCAUGCACAUAUGCGG AUAAUGGUGG	1927
1367	ACCAUUAUGU G ACAUU	1779	AAUGU UGAUGGCAUGCACAUAUGCGG ACAUAAUUGU	1928
1390	CAUAUACAGC G CAAUU	1780	AAUUG UGAUGGCAUGCACAUAUGCGG GCUGUAUAUG	1929
1397	AGCGCAUUU G AACAU	1781	AUGUU UGAUGGCAUGCACAUAUGCGG AAAUUGCGCU	1930
1411	AUACCAUCCU G UUGCG	1782	CGCAA UGAUGGCAUGCACAUAUGCGG AGGAUGGUUU	1931
1414	CCAUCUCUGU G CGUCC	1783	GGACG UGAUGGCAUGCACAUAUGCGG AACAGGAUGG	1932
1425	CGUCCAAU G UAAAG	1784	CUUUA UGAUGGCAUGCACAUAUGCGG AUGUUGGACG	1933
1436	UAAAGAGUU G UCAGC	1785	GCUGA UGAUGGCAUGCACAUAUGCGG AACUUCUUUA	1934
1451	CAGAGGAGAU G ACUAU	1786	AUAGU UGAUGGCAUGCACAUAUGCGG AUCUCCUCUG	1935
1496	UUUAUUUCU G AGCUU	1787	AAGCU UGAUGGCAUGCACAUAUGCGG AGAAAAUAAA	1936
1503	UCUGAGCUUU G UUGGA	1788	UCCAA UGAUGGCAUGCACAUAUGCGG AAAGCUCAGA	1937
1515	UGGAAACAU G AUACC	1789	GGUAU UGAUGGCAUGCACAUAUGCGG AUGUUUUCCA	1938
1532	GAUUUAUUU G CCACA	1790	UGUGG UGAUGGCAUGCACAUAUGCGG AAAUUAUUUC	1939
1539	UUUGCCACAU G UUGUC	1791	GACAA UGAUGGCAUGCACAUAUGCGG AUGUGGCAAA	1940
1542	GCCACAUGUU G UCUGU	1792	ACAGA UGAUGGCAUGCACAUAUGCGG AACAUUGGC	1941
1546	CAUGUUGUCU G UUUUA	1793	UAAAA UGAUGGCAUGCACAUAUGCGG AGACAACAUG	1942
1565	GUGGACCAU G UAAUA	1794	UAUUA UGAUGGCAUGCACAUAUGCGG AUGGGUCCAC	1943
1582	UUUAUCCAU G UUUAA	1795	UAAAA UGAUGGCAUGCACAUAUGCGG AUGGAUAAAA	1944
1624	ACCGUCUAU G UAAUU	1796	AAUUA UGAUGGCAUGCACAUAUGCGG AUUAGACGGU	1945
1637	AUUAAACCAAC G AAAAA	1797	UUUUU UGAUGGCAUGCACAUAUGCGG GUUGGUAUAU	1946
1662	ACUUUUAAU G CUAAC	1798	GUUAG UGAUGGCAUGCACAUAUGCGG AUUUAAAAAGU	1947
1669	AAUGCUAACU G UUUUU	1799	AAAAA UGAUGGCAUGCACAUAUGCGG AGUUAGCAUU	1948
1684	UCCCCUCCU G UCUAG	1800	CUAGA UGAUGGCAUGCACAUAUGCGG AGGAAGGGGA	1949
1696	CUAGGAAAAU G CUUAU	1801	UAUAG UGAUGGCAUGCACAUAUGCGG AUUUUCCUAG	1950
1723	AGUUAGGAAU G ACUUA	1802	UAAGU UGAUGGCAUGCACAUAUGCGG AUUCCUAACU	1951
1737	UAUACGUUUU G UUUUG	1803	CAAAA UGAUGGCAUGCACAUAUGCGG AAAACGUUAU	1952
1742	GUUUUGUUUU G AAUAC	1804	GUAAU UGAUGGCAUGCACAUAUGCGG AAAACAAAAAC	1953
1777	UAUUUAUAUU G CCAUA	1805	UAUGG UGAUGGCAUGCACAUAUGCGG AAUAUAAAAUA	1954

Table 11

1792	AUUCUUACUU G AAUGC	1806	GCAUU UGAUGGCAUGCACUAUGCGG AAGUAAGAAU	1955
1796	UUACUUGAAU G CUUUG	1807	CRAAG UGAUGGCAUGCACUAUGCGG AUUCRAGUAA	1956
1801	UGAAUGCUUU G AAGA	1808	UCAUU UGAUGGCAUGCACUAUGCGG AAAGCAUUA	1957
1805	UGC UUUGAAU G ACUAC	1809	GUAGU UGAUGGCAUGCACUAUGCGG AUUCAAAGCA	1958
1821	AUCCAGUUCU G CACCU	1810	AGGUG UGAUGGCAUGCACUAUGCGG AGAACUGGAU	1959
1839	ACCCUCUGGU G UUGCU	1811	AGCAA UGAUGGCAUGCACUAUGCGG ACCAGAGGGU	1960
1842	CUCUGGUGUU G CUUUU	1812	AAAAG UGAUGGCAUGCACUAUGCGG AACACCAGAG	1961
1924	UUCCAAAAU G ACCAC	1813	GUGGU UGAUGGCAUGCACUAUGCGG GAUUUUGGAA	1962
1961	CUCAGCAAAU G UAAAA	1814	UUUUA UGAUGGCAUGCACUAUGCGG AUUUGCUGAG	1963
1988	AUAACAACU G UCUCU	1815	AGAGA UGAUGGCAUGCACUAUGCGG AGUUUGUUUAU	1964
2067	CAUGGAAACU G AACAA	1816	UUUUU UGAUGGCAUGCACUAUGCGG AGUUUCCAUG	1965
2076	UGAACAAACU G CUCCU	1817	AGGAG UGAUGGCAUGCACUAUGCGG AGGUUGUUUA	1966
2082	ACCUGCUCCU G AAUGA	1818	UCAUU UGAUGGCAUGCACUAUGCGG AGGAGCAGGU	1967
2086	GCUCUGAAU G ACUAC	1819	GUAGU UGAUGGCAUGCACUAUGCGG AUUCAGGAGC	1968
2109	AUAACAACU G AAGGC	1820	GCUUU UGAUGGCAUGCACUAUGCGG AUUUUGUUUAU	1969
2127	AAUAAAGAU G UUCUU	1821	AAGAA UGAUGGCAUGCACUAUGCGG AUUUUUUUUU	1970
2143	UAAACCAAU G AGAAC	1822	GUUCU UGAUGGCAUGCACUAUGCGG AUUGGUUUUA	1971
2193	UCAAGCAGU G UGUAG	1823	CUACA UGAUGGCAUGCACUAUGCGG ACUGCUUUUA	1972
2195	AAAGCAGUGU G UAGAG	1824	CUCUA UGAUGGCAUGCACUAUGCGG ACACUGCUUU	1973
2221	AGCACUAAU G CCCAC	1825	GUGGG UGAUGGCAUGCACUAUGCGG AUUUAGUGCU	1974
2254	AUCUAAAUU G ACACC	1826	GGUGU UGAUGGCAUGCACUAUGCGG AAUUUUUAGAU	1975
2306	CAACACAUU G AAAAG	1827	CUUUU UGAUGGCAUGCACUAUGCGG AAUGUGUUUG	1976
2352	GAGCAGAAU G AAGGA	1828	UCCUU UGAUGGCAUGCACUAUGCGG AGUUCUGCUC	1977
2392	AAAAUCAAU G AAUCC	1829	GGAUU UGAUGGCAUGCACUAUGCGG AUUGAUUUUU	1978
2413	CUGGUUUUU G AAACG	1830	CGUUU UGAUGGCAUGCACUAUGCGG AAAAAACCCAG	1979
2418	UUUUUGAAAC G AUCAA	1831	UUUAU UGAUGGCAUGCACUAUGCGG GUUUCAAAAA	1980
2431	CAACAACU G AUAGA	1832	UUUAU UGAUGGCAUGCACUAUGCGG AUUUUUUUUG	1981
2496	AAUAAAAAU G AUAUA	1833	UUUAU UGAUGGCAUGCACUAUGCGG AUUUUUUUUU	1982

Input Sequence = HSU29607. Cut Site = YG/M or UG/U.

Stem Length = 5/10. Core Sequence = UGAUG GCAUGCACUAUGC GCG

Table 11

Seq1 = HSU29607 (Human methionine aminopeptidase mRNA, complete cds., 2569 bp)		

Table 12

Table 12: Anti Human MetAP-2 HH, NCH, and G-Cleaver Ribozymes

Alias	Ribozyme Sequence	Seq. ID Nos.	Substrate Seq.	Seq. ID Nos.
HH				
MAP2-11	CCCGAGA CUGAUGAGGCCGCUUAGGCCGAA AGACGAG	1983	CUCGUCU C UCUCGGG	2001
MAP2-15	GUUGCCC CUGAUGAGGCCGCUUAGGCCGAA AGAGAGA	1984	UCUCUCU C GGGCAAC	2002
MAP2-464	GUUCUCC CUGAUGAGGCCGCUUAGGCCGAA AGCAGCU	1985	AGCUGCU U GGAGAAC	2003
MAP2-911	UCACACA CUGAUGAGGCCGCUUAGGCCGAA ACGAACA	1986	UGUUCGU C UGUGUGA	2004
MAP2-1290	UUCUCCC CUGAUGAGGCCGCUUAGGCCGAA AGCGAUC	1987	GAUCGCU U GGAGAGAA	2005
MAP2-1342	GAUCUAC CUGAUGAGGCCGCUUAGGCCGAA AUGCCCA	1988	UGGGCAU U GUAGAUC	2006
MAP2-1479	AGGUGU CUGAUGAGGCCGCUUAGGCCGAA AGGUGGC	1989	GCCACCU C AACACCU	2007
MAP2-1646	GUCCGGA CUGAUGAGGCCGCUUAGGCCGAA AGCUUUU	1990	AAAAGCU U UCCGGAC	2008
MAP2-1819	AGGUGCA CUGAUGAGGCCGCUUAGGCCGAA AACUGGA	1991	UCCAGU C UGCACCU	2009
MAP2-2262	GUAGU CUGAUGAGGCCGCUUAGGCCGAA AGGUGU	1992	ACACCU A ACAUCAC	2010
MAP2-10	CCGAGAG CUGAUGAGGCCGCUUAGGCCGAA GACGAGG	1993	CCUGUC U CUCUGGG	2011
NCH				
MAP2-369	AACUGAG CUGAUGAGGCCGCUUAGGCCGAA IAGGGUC	1994	GACCCUC C CUCAGUU	2012
MAP2-370	GAACUGA CUGAUGAGGCCGCUUAGGCCGAA IGAGGGU	1995	ACCCUCC C UCAGUUC	2013
MAP2-1901	GUUGUGU CUGAUGAGGCCGCUUAGGCCGAA IUGCUGA	1996	UCAGCAC C ACACAAC	2014
MAP2-1906	UAGGUGU CUGAUGAGGCCGCUUAGGCCGAA IUGUGGU	1997	ACCACAC A ACACCUA	2015
G-Cleaver				
MAP2-1821	AGGUG UGAUGGCAUGCACAUGCGCG AGAACUGGAU	1998	AUCCAGUUCU G CACCU	2016
MAP2-2076	AGGAG UGAUGGCAUGCACAUGCGCG AGGUUGUUA	1999	UGAACAAACCU G CUCCU	2017
MAP2-2086	GUAGU UGAUGGCAUGCACAUGCGCG AUUCAGGAGC	2000	GCUCUGAAU G ACUAC	2018

Table 13

Table 13: Human telomerase reverse transcriptase (TERT) Hammerhead Ribozyme and Target Sequence

nt. Position	Ribozyme Sequence	Seq ID Nos.	Substrate Sequence	Seq ID Nos.
13	CGCAGCAG CUGAUGAG X CGAA ACGCAGCG		CGCTGCGT C CTGCTGCG	
68	GCAGCGGG CUGAUGAG X CGAA AGCGCGCG		CGCGCGCT C CCCGCTGC	
90	GCAGCAGG CUGAUGAG X CGAA AGCGCACG		CGTGCGCT C CCTGCTGC	
108	CCUCGCGG CUGAUGAG X CGAA AGUGGCUG		CAGCCACT A CCGCAGAG	
135	GCCGCACG CUGAUGAG X CGAA ACGUGGCC		GGCCACGT T CGTGCGGC	
136	CGCCGCAC CUGAUGAG X CGAA AACGUGGC		GCCACGTT C GTGCGGCG	
194	CGCGCGGA CUGAUGAG X CGAA AGCCGCCG		CGGCGGCT T TCCGCGCG	
195	GCGCGCGG CUGAUGAG X CGAA AAGCCGCC		GGCGGCTT T CCGCGCGC	
196	AGCGCGCG CUGAUGAG X CGAA AAAGCCGC		GCGGCTTT C CGCGCGCT	
264	GGCGGAAG CUGAUGAG X CGAA AGGGGGCG		CGCCCCCT C CTTCCGCC	
267	CCUGGCGG CUGAUGAG X CGAA AGGAGGGG		CCCCTCCT T CCGCCAGG	
268	ACCUGGCG CUGAUGAG X CGAA AAGGAGGG		CCCTCCTT C CGCCAGGT	
279	UCAGGCAG CUGAUGAG X CGAA ACACCUGG		CCAGGTGT C CTGCCTGA	
351	CGAAGCCG CUGAUGAG X CGAA AGGCCAGC		GCTGGCCT T CGGCTTCG	
352	GCGAAGCC CUGAUGAG X CGAA AAGGCCAG		CTGGCCTT C GGCTTCGC	
357	GCAGCGCG CUGAUGAG X CGAA AGCCGAAG		CTTCGGCT T CGCGCTGC	
358	AGCAGCGC CUGAUGAG X CGAA AAGCCGAA		TTCGGCTT C GCGCTGCT	
399	UGGUGGUG CUGAUGAG X CGAA AGGCCUCG		CGAGGCCT T CACCACCA	
400	CUGGUGGU CUGAUGAG X CGAA AAGGCCUC		GAGGCCTT C ACCACCAG	
420	UGGGCAGG CUGAUGAG X CGAA AGCUGCGC		GCGCAGCT A CCTGCCCA	
505	AGCAGGUG CUGAUGAG X CGAA ACCAGCAC		GTGCTGGT T CACCTGCT	
506	CAGCAGGU CUGAUGAG X CGAA AACCAGCA		TGCTGGTT C ACCTGCTG	
529	AGCACAAA CUGAUGAG X CGAA AGCGCGCA		TGCGCGCT C TTTGTGCT	
531	CCAGCACA CUGAUGAG X CGAA AGAGCGCG		CGCGCTCT T TGTGCTGG	
532	ACCAGCAC CUGAUGAG X CGAA AAGAGCGC		GCGCTCTT T GTGCTGGT	
545	GCAGCUGG CUGAUGAG X CGAA AGCCACCA		TGGTGGCT C CCAGCTGC	
558	ACACCUGG CUGAUGAG X CGAA AGGCGCAG		CTGCGCCT A CCAGGTGT	
582	CGAGCUGG CUGAUGAG X CGAA ACAGCGGC		GCCGCTGT A CCAGCTCG	
589	GCAGCGCC CUGAUGAG X CGAA AGCUGGUA		TACCAGCT C GCGCTGCT	
602	CCGGGCCU CUGAUGAG X CGAA AGUGGCAG		CTGCCACT C AGGCCCGG	
626	GGGUCCAC CUGAUGAG X CGAA AGCGUGUG		CACACGCT A GTGGACCC	
644	GCAUCCCA CUGAUGAG X CGAA ACGCCUUC		GAAGGCGT C TGGGATGC	
671	CCUGACGC CUGAUGAG X CGAA AUGGUUCC		GGAACCAT A GCGTCAGG	
676	GCCUCCCU CUGAUGAG X CGAA ACGCUAUG		CATAGCGT C AGGGAGGC	
691	CCCAGGGG CUGAUGAG X CGAA ACCCCGGC		GCCGGGGT C CCCCTGGG	
749	CAACGGCA CUGAUGAG X CGAA ACUUCGGC		GCCGAAGT C TGCCGTGT	
756	UCUUGGGC CUGAUGAG X CGAA ACGGCAGA		TCTGCCGT T GCCCAAGA	
808	CCCUGCCC CUGAUGAG X CGAA ACGGGCGU		ACGCCCGT T GGGCAGGG	
819	GGGCCCAG CUGAUGAG X CGAA ACCCCUGC		GCAGGGGT C CTGGGCCC	
863	CACACAGA CUGAUGAG X CGAA ACCACGGU		ACCGTGGT T TCTGTGTG	
864	CCACACAG CUGAUGAG X CGAA AACCACGG		CCGTGGTT T CTGTGTGG	
865	ACCACACA CUGAUGAG X CGAA AAACCACG		CGTGGTTT C TGTGTGGT	
876	UGGCAGGU CUGAUGAG X CGAA ACACCACA		TGTGGTGT C ACCTGCCA	

Table 13

906	CCUCCAAA CUGAUGAG X CGAA AGGUGGCU	AGCCACCT C TTTGGAGG
908	ACCCUCCA CUGAUGAG X CGAA AGAGGUGG	CCACCTCT T TGGAGGGT
909	CACCCUCC CUGAUGAG X CGAA AAGAGGUG	CACCTCTT T GGAGGGTG
922	GUGCCAGA CUGAUGAG X CGAA AGCGCACC	GGTGCCT C TCTGGCAC
924	GCGUGCCA CUGAUGAG X CGAA AGAGCGCA	TGCGCTCT C TGGCACGC
939	AUGGGUGG CUGAUGAG X CGAA AGUGGCGC	GCGCCACT C CCACCCAT
948	GGCCCACG CUGAUGAG X CGAA AUGGGUGG	CCACCCAT C CGTGGGCC
981	GCGAUGUG CUGAUGAG X CGAA AUGGGGGG	CCCCCAT C CACATCGC
987	GUGGCCGC CUGAUGAG X CGAA AUGUGGAU	ATCCACAT C GCGGCCAC
1001	GUCCCAGG CUGAUGAG X CGAA ACGUGGUG	CACCACGT C CCTGGGAC
1016	CGGGGGAC CUGAUGAG X CGAA AGCGGUGU	ACACGCCT T GTCCCCCG
1019	CACCGGGG CUGAUGAG X CGAA ACAAGGCG	CGCCTTGT C CCCCAGTG
1029	UCUCGGCG CUGAUGAG X CGAA ACACCGGG	CCCGTGT A CGCCGAGA
1047	AGUAGAGG CUGAUGAG X CGAA AGUGCUUG	CAAGCACT T CCTCTACT
1048	GAGUAGAG CUGAUGAG X CGAA AAGUGCUU	AAGCACTT C CTCTACTC
1051	GAGGAGUA CUGAUGAG X CGAA AGGAAGUG	CACTTCCT C TACTCCTC
1053	CUGAGGAG CUGAUGAG X CGAA AGAGGAAG	CTTCCTCT A CTCCTCAG
1056	CGCCUGAG CUGAUGAG X CGAA AGUAGAGG	CCTCTACT C CTCAGGCG
1059	UGUCGCCU CUGAUGAG X CGAA AGGAGUAG	CTACTCCT C AGGCGACA
1086	GUAGGAAG CUGAUGAG X CGAA AGGGCCGC	GCGGCCCT C TTCTCTAC
1089	UGAGUAGG CUGAUGAG X CGAA AGGAGGGC	GCCCTCCT T CCTACTCA
1090	CUGAGUAG CUGAUGAG X CGAA AAGGAGGG	CCCTCCTT C CTACTCAG
1093	GAGCUGAG CUGAUGAG X CGAA AGGAAGGA	TCCTTCCT A CTCAGCTC
1096	AGAGAGCU CUGAUGAG X CGAA AGUAGGAA	TTCCTACT C AGCTCTCT
1101	GCCUCAGA CUGAUGAG X CGAA AGCUGAGU	ACTCAGCT C TCTGAGGC
1103	GGGCCUCA CUGAUGAG X CGAA AGAGCUGA	TCAGCTCT C TGAGGCCC
1127	GAGCCUCC CUGAUGAG X CGAA AGCGCCAG	CTGGCGCT C GGAGGCTC
1135	GUCUCCAC CUGAUGAG X CGAA AGCCUCCG	CGGAGGCT C GTGGAGAC
1147	CCCAGAAA CUGAUGAG X CGAA AUGGUCUC	GAGACCAT C TTTCTGGG
1149	AAACCAGA CUGAUGAG X CGAA AGAUGGUC	GACCATCT T TCTGGGTT
1150	GAACCAG CUGAUGAG X CGAA AAGAUGGU	ACCATCTT T CTGGGTTT
1151	GGAACCCA CUGAUGAG X CGAA AAAGAUGG	CCATCTTT C TGGGTTCC
1157	GGGCCUGG CUGAUGAG X CGAA ACCCAGAA	TTCTGGGT T CCAGGCCC
1158	AGGGCCUG CUGAUGAG X CGAA AACCCAGA	TCTGGGTT C CAGGCCCT
1181	CCUGCGGG CUGAUGAG X CGAA AGUCCUG	CAGGGACT C CCCGAGG
1191	GGCGGGGC CUGAUGAG X CGAA ACCUGCGG	CCGAGGT T GCCCCGCC
1212	UUUGCCAG CUGAUGAG X CGAA AGCGCUGG	CCAGCGCT A CTGGCAAA
1233	GCUCCAGA CUGAUGAG X CGAA ACAGGGGC	GCCCTGT T TCTGGAGC
1234	AGCUCCAG CUGAUGAG X CGAA AACAGGGG	CCCCTGTT T CTGGAGCT
1235	CAGCUCCA CUGAUGAG X CGAA AACAGGGG	CCCTGTTT C TGGAGCTG
1246	UGGUUCCC CUGAUGAG X CGAA AGCAGCUC	GAGCTGCT T GGGAACCA
1269	GCACCCCG CUGAUGAG X CGAA AGGGGCAC	GTGCCCCCT A CGGGGTGC
1279	GUCUUGAG CUGAUGAG X CGAA AGCACCCC	GGGGTGCT C CTCAAGAC
1282	UGCGUCUU CUGAUGAG X CGAA AGGAGCAC	GTGTCCT C AAGACGCA
1312	GCUGGGGU CUGAUGAG X CGAA ACCGCAGC	GCTGCGGT C ACCCCAGC
1330	CGGGCACA CUGAUGAG X CGAA ACACCGGC	GCCGGTGT C TGTGCCCG
1356	CCGCCACA CUGAUGAG X CGAA AGCCUGG	CCAGGGCT C TGTGGCGG

Table 13

1394	CACCAGGC CUGAUGAG X CGAA ACGGGGGU	ACCCCCGT C GCCTGGTG
1411	UGCUGGCG CUGAUGAG X CGAA AGCAGCUG	CAGCTGCT C CGCCAGCA
1440	CGAAGCCG CUGAUGAG X CGAA ACACCUGC	GCAGGTGT A CGGCTTCG
1446	CCCGCACG CUGAUGAG X CGAA AGCCGUAC	GTACGGCT T CGTGGCGG
1447	GCCCGCAC CUGAUGAG X CGAA AAGCCGUA	TACGGCTT C GTGGGGGC
1486	GAGCCCCA CUGAUGAG X CGAA AGGCCUGG	CCAGGCCT C TGGGGCTC
1494	UGUGCCUG CUGAUGAG X CGAA AGCCCCAG	CTGGGGCT C CAGGCACA
1515	UCCUGAGG CUGAUGAG X CGAA AGCGGCGU	ACGCCGCT T CCTCAGGA
1516	UUCUGAG CUGAUGAG X CGAA AAGCGGCG	CGCCGCTT C CTCAGGAA
1519	GUGUCCU CUGAUGAG X CGAA AGGAAGCG	CGCTTCCT C AGGAACAC
1536	GGGAGAUG CUGAUGAG X CGAA ACUUCUUG	CAAGAAGT T CATCTCCC
1537	AGGGAGAU CUGAUGAG X CGAA AACUUCUU	AAGAAGTT C ATCTCCCT
1540	CCCAGGGA CUGAUGAG X CGAA AUGAACUU	AAGTTCAT C TCCCTGGG
1542	UCCCCAGG CUGAUGAG X CGAA AGAUGAAC	GTTCATCT C CCTGGGGA
1564	UGCAGCGA CUGAUGAG X CGAA AGCUUGGC	GCCAAGCT C TCGCTGCA
1566	CCUGCAGC CUGAUGAG X CGAA AGAGCUUG	CAAGCTCT C GCTGCAGG
1610	GCGCAGCC CUGAUGAG X CGAA AGCGCAGU	ACTGCGCT T GGCTGCGC
1633	ACACAGCC CUGAUGAG X CGAA ACCCCUGG	CCAGGGGT T GGCTGTGT
1642	GCGGCCGG CUGAUGAG X CGAA ACACAGCC	GGCTGTGT T CCGGCCGC
1643	UGCGGCCG CUGAUGAG X CGAA AACACAGC	GCTGTGTT C CGGCCGCA
1661	CUCACGCA CUGAUGAG X CGAA ACGGUGCU	AGCACCGT C TCGTGAG
1675	UUGGCCAG CUGAUGAG X CGAA AUCUCCUC	GAGGAGAT C CTGGCCAA
1686	AGUGCAGG CUGAUGAG X CGAA ACUUGGCC	GGCCAAGT T CCTGCACT
1687	CAGUGCAG CUGAUGAG X CGAA AACUUGGC	GCCAAGTT C CTGCACTG
1710	CGACGACG CUGAUGAG X CGAA ACACACUC	GAGTGTGT A CGTCGTCG
1714	AGCUCGAC CUGAUGAG X CGAA ACGUACAC	GTGTACGT C GTCGAGCT
1717	AGCAGCUC CUGAUGAG X CGAA ACGACGUA	TACGTCGT C GAGCTGCT
1726	AAAGACCU CUGAUGAG X CGAA AGCAGCUC	GAGCTGCT C AGGTCTTT
1731	AAAAGAAA CUGAUGAG X CGAA ACCUGAGC	GCTCAGGT C TTTCTTTT
1733	AUAAAAGA CUGAUGAG X CGAA AGACCUGA	TCAGGTCT T TCTTTTAT
1734	CAUAAAAG CUGAUGAG X CGAA AAGACCUG	CAGGTCTT T CTTTATATG
1735	ACAUAAAA CUGAUGAG X CGAA AAAGACCU	AGGTCTTT C TTTTATGT
1737	UGACAUAA CUGAUGAG X CGAA AGAAAGAC	GTCTTTCT T TTATGTCA
1738	GUGACAU CUGAUGAG X CGAA AAGAAAGA	TCTTTCTT T TATGTCAC
1739	CGUGACAU CUGAUGAG X CGAA AAAGAAAG	CTTTCTTT T ATGTCACG
1740	CCGUGACA CUGAUGAG X CGAA AAAAGAAA	TTTCTTTT A TGTCACGG
1744	GUCUCCGU CUGAUGAG X CGAA ACAUAAAA	TTTTATGT C ACGGAGAC
1758	UCUUUUGA CUGAUGAG X CGAA ACGUGGUC	GACCACGT T TCAAAAGA
1759	UUCUUUUG CUGAUGAG X CGAA AACGUGGU	ACCACGTT T CAAAAGAA
1760	GUUCUUUU CUGAUGAG X CGAA AAACGUGG	CCACGTTT C AAAAGAAC
1774	UAGAAAAA CUGAUGAG X CGAA AGCCUGUU	AACAGGCT C TTTTCTA
1776	GGUAGAAA CUGAUGAG X CGAA AGAGCCUG	CAGGCTCT T TTTCTACC
1777	CGGUAGAA CUGAUGAG X CGAA AAGAGCCU	AGGCTCTT T TTCTACCG
1778	CCGGUAGA CUGAUGAG X CGAA AAAGAGCC	GGCTCTTT T TCTACCGG
1779	UCCGGUAG CUGAUGAG X CGAA AAAAGAGC	GCTCTTTT T CTACCGGA
1780	UUCCGGUA CUGAUGAG X CGAA AAAAAGAG	CTCTTTT C TACCGGAA
1782	UCUCCGG CUGAUGAG X CGAA AGAAAAAG	CTTTTCT A CCGGAAGA

Table 13

1795	UUGCUGCA CUGAUGAG X CGAA ACACUCUU	AAGAGTGT C TGGAGCAA
1806	UGCUGGC CUGAUGAG X CGAA ACUUGCUC	GAGCAAGT T GCAAAGCA
1816	CUGAUCC CUGAUGAG X CGAA AUGCUUUG	CAAAGCAT T GGAATCAG
1822	UGCUGUC CUGAUGAG X CGAA AUUCCAAU	ATTGGAAT C AGACAGCA
1833	CCCUCUC CUGAUGAG X CGAA AGUGCUGU	ACAGCACT T GAAGAGGG
1860	CUGCUCC CUGAUGAG X CGAA ACAGCUCC	GGAGCTGT C GGAAGCAG
1873	UGCUGCCU CUGAUGAG X CGAA ACCUCUGC	GCAGAGGT C AGGCAGCA
1883	GGCUUCC CUGAUGAG X CGAA AUGCUGCC	GGCAGCAT C GGAAGCC
1911	GGAGUCUG CUGAUGAG X CGAA ACGUCAGC	GCTGACGT C CAGACTCC
1918	AUGAAGCG CUGAUGAG X CGAA AGUCUGGA	TCCAGACT C CGCTTCAT
1923	UGGGGAUG CUGAUGAG X CGAA AGCGGAGU	ACTCCGCT T CATCCCCA
1924	UUGGGGAU CUGAUGAG X CGAA AAGCGGAG	CTCCGCTT C ATCCCCAA
1927	GGCUUGGG CUGAUGAG X CGAA AUGAAGCG	CGCTTCAT C CCCAAGCC
1954	AUGUUCAC CUGAUGAG X CGAA AUCGGCCG	CGGCCGAT T GTGAACAT
1968	CCACGACG CUGAUGAG X CGAA AGUCCAUG	CATGGACT A CGTCGTGG
1972	GCUCCAC CUGAUGAG X CGAA ACGUAGUC	GACTACGT C GTGGGAGC
1989	CUCUGCG CUGAUGAG X CGAA ACGUUCUG	CAGAACGT T CCGCAGAG
1990	UCUCUGCG CUGAUGAG X CGAA AACGUUCU	AGAACGTT C CGCAGAGA
2015	CGAGGUGA CUGAUGAG X CGAA ACGCUCGG	CCGAGCGT C TCACCTCG
2017	CUCGAGGU CUGAUGAG X CGAA AGACGCUC	GAGCGTCT C ACCTCGAG
2022	UCACCCUC CUGAUGAG X CGAA AGGUGAGA	TCTCACCT C GAGGGTGA
2040	GCACGCG CUGAUGAG X CGAA ACAGUGCC	GGCACTGT T CAGCGTGC
2041	AGCACGCU CUGAUGAG X CGAA AACAGUGC	GCACTGTT C AGCGTGCT
2050	UCGUAGUU CUGAUGAG X CGAA AGCACGCU	AGCGTGCT C AACTACGA
2055	CCCGCUCG CUGAUGAG X CGAA AGUUGAGC	GCTCAACT A CGAGCGGG
2080	GCGCCAG CUGAUGAG X CGAA AGGCCGGG	CCCGGCCT C CTGGGCGC
2091	CCAGCACA CUGAUGAG X CGAA AGGCGCCC	GGGCGCCT C TGTGCTGG
2111	CCUGUGGA CUGAUGAG X CGAA AUCGUCCA	TGGACGAT A TCCACAGG
2113	GCCUGUG CUGAUGAG X CGAA AUAUCGUC	GACGATAT C CACAGGGC
2133	GCAGCACG CUGAUGAG X CGAA AGGUGCGC	GCGCACCT T CGTGCTGC
2134	CGCAGCAC CUGAUGAG X CGAA AAGGUGCG	CGCACCTT C GTGCTGCG
2175	UGACAAAG CUGAUGAG X CGAA ACAGCUCA	TGAGCTGT A CTTTGTCA
2178	CCUUGACA CUGAUGAG X CGAA AGUACAGC	GCTGTACT T TGTCAAGG
2179	ACCUUGAC CUGAUGAG X CGAA AAGUACAG	CTGTACTT T GTCAAGGT
2182	UCCACCUU CUGAUGAG X CGAA ACAAAGUA	TACTTTGT C AAGGTGGA
2205	UGGUGUCG CUGAUGAG X CGAA ACGCGCCC	GGGCGCGT A CGACACCA
2215	UCCUGGGG CUGAUGAG X CGAA AUGGUGUC	GACACCAT C CCCAGGA
2230	ACCUCCGU CUGAUGAG X CGAA AGCCUGUC	GACAGGCT C ACGGAGGT
2239	CUGGCGAU CUGAUGAG X CGAA ACCUCCGU	ACGGAGGT C ATGCCAG
2242	AUGCUGGC CUGAUGAG X CGAA AUGACCUC	GAGGTCAT C GCCAGCAT
2251	GGUUGAU CUGAUGAG X CGAA AUGCUGGC	GCCAGCAT C ATCAAACC
2254	UGGGGUUU CUGAUGAG X CGAA AUGAUGCU	AGCATCAT C AAACCCCA
2271	GCACGCAG CUGAUGAG X CGAA ACGUGUUC	GAACACGT A CTGCGTGC
2282	GGCAUACC CUGAUGAG X CGAA ACGCACGC	GCGTGCGT C GGTATGCC
2286	CCACGGCA CUGAUGAG X CGAA ACCGACGC	GCGTCGGT A TGCCGTGG
2296	GCCUUCUG CUGAUGAG X CGAA ACCACGGC	GCCGTGGT C CAGAAGGC
2320	GCCUUGCG CUGAUGAG X CGAA ACGUGCCC	GGGCACGT C CGCAAGGC

Table 13

2331	GGCUCUUG CUGAUGAG X CGAA AGGCCUUG	CAAGGCCT T CAAGAGCC
2332	UGGCUCUU CUGAUGAG X CGAA AAGGCCUU	AAGGCCTT C AAGAGCCA
2344	AAGGUAGA CUGAUGAG X CGAA ACGUGGCU	AGCCACGT C TCTACCTT
2346	UCAAGGUA CUGAUGAG X CGAA AGACGUGG	CCACGTCT C TACCTTGA
2348	UGUCAAGG CUGAUGAG X CGAA AGAGACGU	ACGTCTCT A CCTTGACA
2352	GGUCUGUC CUGAUGAG X CGAA AGGUAGAG	CTCTACCT T GACAGACC
2362	UACGGCUG CUGAUGAG X CGAA AGGUCUGU	ACAGACCT C CAGCCGTA
2370	GUCGCAUG CUGAUGAG X CGAA ACGGCUUG	CCAGCCGT A CATGCGAC
2382	GAGCCACG CUGAUGAG X CGAA ACUGUCGC	GCGACAGT T CGTGGCTC
2383	UGAGCCAC CUGAUGAG X CGAA AACUGUCG	CGACAGTT C GTGGCTCA
2390	CUGCAGGU CUGAUGAG X CGAA AGCCACGA	TCGTGGCT C ACCTGCAG
2425	UCGAUGAC CUGAUGAG X CGAA ACGGCAUC	GATGCCGT C GTCATCGA
2428	UGCUCGAU CUGAUGAG X CGAA ACGACGGC	GCCGTCGT C ATCGAGCA
2431	CUCUGCUC CUGAUGAG X CGAA AUGACGAC	GTCGTCAT C GAGCAGAG
2442	UCAGGGAG CUGAUGAG X CGAA AGCUCUGC	GCAGAGCT C CTCCTGA
2445	CAUUCAGG CUGAUGAG X CGAA AGGAGCUC	GAGCTCCT C CCTGAATG
2470	ACGUCGAA CUGAUGAG X CGAA AGGCCACU	AGTGGCCT C TTCGACGT
2472	AGACGUCG CUGAUGAG X CGAA AGAGGCCA	TGGCCTCT T CGACGTCT
2473	AAGACGUC CUGAUGAG X CGAA AAGAGGCC	GGCCTCTT C GACGTCTT
2479	CGUAGGAA CUGAUGAG X CGAA ACGUCGAA	TTCGACGT C TTCCTACG
2481	AGCGUAGG CUGAUGAG X CGAA AGACGUCG	CGACGTCT T CCTACGCT
2482	AAGCGUAG CUGAUGAG X CGAA AAGACGUC	GACGTCTT C CTACGCTT
2485	AUGAAGCG CUGAUGAG X CGAA AGGAAGAC	GTCTTCCT A CGCTTCAT
2490	GGCACAU CUGAUGAG X CGAA AGCGUAGG	CCTACGCT T CATGTGCC
2491	UGGCACAU CUGAUGAG X CGAA AAGCGUAG	CTACGCTT C ATGTGCCA
2515	UUGCCCUU CUGAUGAG X CGAA AUGCGCAC	GTGCGCAT C AGGGGCAA
2526	GGACGUAG CUGAUGAG X CGAA ACUUGCCC	GGGCAAGT C CTACGTCC
2529	ACUGGACG CUGAUGAG X CGAA AGGACUUG	CAAGTCCT A CGTCCAGT
2533	UGGCACUG CUGAUGAG X CGAA ACGUAGGA	TCCTACGT C CAGTGCCA
2548	CCUGCGG CUGAUGAG X CGAA AUCCCUUG	CAGGGGAT C CCGCAGGG
2559	AGAGGAUG CUGAUGAG X CGAA AGCCUUGC	GCAGGGCT C CATCCTCT
2563	GUGGAGAG CUGAUGAG X CGAA AUGGAGCC	GGCTCCAT C CTCTCCAC
2566	AGCGUGGA CUGAUGAG X CGAA AGGAUGGA	TCCATCCT C TCCACGCT
2568	GCAGCGUG CUGAUGAG X CGAA AGAGGAUG	CATCCTCT C CACGCTGC
2578	AGGCUGCA CUGAUGAG X CGAA AGCAGCGU	ACGCTGCT C TGCAGCCT
2592	UGUCGCCG CUGAUGAG X CGAA AGCACAGG	CCTGTGCT A CGGCGACA
2616	UCCCCGCA CUGAUGAG X CGAA ACAGCUUG	CAAGCTGT T TGCGGGGA
2617	AUCCCCGC CUGAUGAG X CGAA AACAGCUU	AAGCTGTT T GCGGGGAT
2626	UCCCGCCG CUGAUGAG X CGAA AUCCCCGC	GCGGGGAT T CGGCGGGA
2627	GUCCCCGC CUGAUGAG X CGAA AAUCCCCG	CGGGGATT C GCGGGGAC
2644	AAACGCAG CUGAUGAG X CGAA AGCAGCCC	GGGCTGCT C CTGCGTTT
2651	AUCCACCA CUGAUGAG X CGAA ACGCAGGA	TCCTGCGT T TGGTGGAT
2652	CAUCCACC CUGAUGAG X CGAA AACGCAGG	CCTGCGTT T GGTGGATG
2663	CAACAAGA CUGAUGAG X CGAA AUCAUCCA	TGGATGAT T TCTTGTG
2664	CCAACAAG CUGAUGAG X CGAA AAUCAUCC	GGATGATT T CTTGTGTTG
2665	ACCAACAA CUGAUGAG X CGAA AAUCAUCC	GATGATTT C TTGTTGGT
2667	UCACCAAC CUGAUGAG X CGAA AGAAAUCA	TGATTTCT T GTTGGTGA

Table 13

2670	GUGUCACC CUGAUGAG X CGAA ACAAGAAA	TTTCTTGT T GGTGACAC
2681	GGUGAGGU CUGAUGAG X CGAA AGGUGUCA	TGACACCT C ACCTCACC
2686	GCGUGGGU CUGAUGAG X CGAA AGGUGAGG	CCTCACCT C ACCCACGC
2703	UCCUGAGG CUGAUGAG X CGAA AGGUUUUC	GAAAACCT T CCTCAGGA
2704	GUCCUGAG CUGAUGAG X CGAA AAGGUUUU	AAAACCTT C CTCAGGAC
2707	AGGGUCCU CUGAUGAG X CGAA AGGAAGGU	ACCTTCCT C AGGACCTT
2719	ACACCUCG CUGAUGAG X CGAA ACCAGGGU	ACCCTGGT C CGAGGTGT
2728	UACUCAGG CUGAUGAG X CGAA ACACCUCG	CGAGGTGT C CCTGAGTA
2736	CGCAGCCA CUGAUGAG X CGAA ACUCAGGG	CCCTGAGT A TGGCTGCG
2754	UCUCCGC CUGAUGAG X CGAA AGUUCACC	GGTGAAC T GCGGAAGA
2775	CUACAGGG CUGAUGAG X CGAA AGUUCACC	GGTGAAC T CCCTGTAG
2776	UCUACAGG CUGAUGAG X CGAA AAGUUCAC	GTGAACCT C CCTGTAGA
2782	UCGUCUUC CUGAUGAG X CGAA ACAGGGAA	TTCCCTGT A GAAGACGA
2810	CUGAACAA CUGAUGAG X CGAA AGCCGUGC	GCACGGCT T TTGTTTCA
2811	UCUGAACA CUGAUGAG X CGAA AAGCCGUG	CACGGCTT T TGTTCAGA
2812	AUCUGAAC CUGAUGAG X CGAA AAAGCCGU	ACGGCTTT T GTTCAGAT
2815	GGCAUCUG CUGAUGAG X CGAA ACAAAGC	GCTTTTGT T CAGATGCC
2816	CGGCAUCU CUGAUGAG X CGAA AACAAAAG	CTTTTGT C AGATGCCG
2836	CAGGGGAA CUGAUGAG X CGAA AGGCCGUG	CACGGCCT A TTCCCCTG
2838	ACCAGGGG CUGAUGAG X CGAA AUAGGCCG	CGGCCTAT T CCCCTGGT
2839	CACCAGGG CUGAUGAG X CGAA AAUAGGCC	GGCCTATT C CCCTGGTG
2864	GGUCCGGG CUGAUGAG X CGAA AUCCAGCA	TGCTGGAT A CCCGGACC
2892	AGCUGGAG CUGAUGAG X CGAA AGUCGCUC	GAGCGACT A CTCCAGCT
2895	CAUAGCUG CUGAUGAG X CGAA AGUAGUCG	CGACTACT C CAGCTATG
2901	UCCGGGCA CUGAUGAG X CGAA AGCUGGAG	CTCCAGCT A TGCCCGGA
2913	CUCUGAUG CUGAUGAG X CGAA AGGUCCGG	CCGGACCT C CATCAGAG
2917	CUGGCUCU CUGAUGAG X CGAA AUGGAGGU	ACCTCCAT C AGAGCCAG
2927	GAAGGUGA CUGAUGAG X CGAA ACUGGCUC	GAGCCAGT C TCACCTTC
2929	UUGAAGGU CUGAUGAG X CGAA AGACUGGC	GCCAGTCT C ACCTTCAA
2934	CGCGGUUG CUGAUGAG X CGAA AGGUGAGA	TCTCACCT T CAACCGCG
2935	CCGCGGUU CUGAUGAG X CGAA AAGGUGAG	CTCACCTT C AACCGCGG
2946	CAGCCUUG CUGAUGAG X CGAA AGCCGCGG	CCGCGGCT T CAAGGCTG
2947	CCAGCCUU CUGAUGAG X CGAA AAGCCGCG	CGCGGCTT C AAGGCTGG
2969	GAGUUUGC CUGAUGAG X CGAA ACGCAUGU	ACATGCGT C GCAAACCTC
2977	ACCCCAA CUGAUGAG X CGAA AGUUUGCG	CGCAAACCT C TTTGGGGT
2979	AGACCCCA CUGAUGAG X CGAA AGAGUUUG	CAAACCTT T TGGGGTCT
2980	AAGACCCC CUGAUGAG X CGAA AAGAGUUU	AAACTCTT T GGGGTCTT
2986	AGCCGCAA CUGAUGAG X CGAA ACCCCAAA	TTTGGGGT C TTGCGGCT
2988	UCAGCCGC CUGAUGAG X CGAA AGACCCCA	TGGGGTCT T GCGGCTGA
3002	CAGGCUGU CUGAUGAG X CGAA ACACUUCA	TGAAGTGT C ACAGCCTG
3012	AAUCCAGA CUGAUGAG X CGAA ACAGGCUG	CAGCCTGT T TCTGGATT
3013	AAAUCCAG CUGAUGAG X CGAA AACAGGCU	AGCCTGTT T CTGGATTT
3014	CAAAUCCA CUGAUGAG X CGAA AAACAGGC	GCCTGTTT C TGGATTTG
3020	CACCUGCA CUGAUGAG X CGAA AUCCAGAA	TTCTGGAT T TGCAGGTG
3021	UCACCUGC CUGAUGAG X CGAA AAUCCAGA	TCTGGATT T GCAGGTGA
3037	ACCGUCUG CUGAUGAG X CGAA AGGCUGUU	AACAGCCT C CAGACGGT
3058	AUCUUGUA CUGAUGAG X CGAA AUGUUGGU	ACCAACAT C TACAAGAT

Table 13

3060	GGAUCUUG CUGAUGAG X CGAA AGAUGUUG	CAACATCT A CAAGATCC
3067	AGCAGGAG CUGAUGAG X CGAA AUCUUGUA	TACAAGAT C CTCCTGCT
3070	UGCAGCAG CUGAUGAG X CGAA AGGAUCUU	AAGATCCT C CTGCTGCA
3084	GAAACCUG CUGAUGAG X CGAA ACGCCUGC	GCAGGCGT A CAGGTTTC
3090	AUGCGUGA CUGAUGAG X CGAA ACCUGUAC	GTACAGGT T TCACGCAT
3091	CAUGCGUG CUGAUGAG X CGAA AACCUGUA	TACAGGTT T CACGCATG
3092	ACAUGCGU CUGAUGAG X CGAA AAACCUGU	ACAGGTTT C ACGCATGT
3112	UGAAAUGG CUGAUGAG X CGAA AGCUGCAG	CTGCAGCT C CCATTTC A
3117	GCUGAUGA CUGAUGAG X CGAA AUGGGAGC	GCTCCCAT T TCATCAGC
3118	UGCUGAUG CUGAUGAG X CGAA AAUGGGAG	CTCCCAT T CATCAGCA
3119	UUGCUGAU CUGAUGAG X CGAA AAAUGGGA	TCCCATT C ATCAGCAA
3122	AACUUGCU CUGAUGAG X CGAA AUGAAAUG	CATTTCAT C AGCAAGTT
3130	UUCUCCA CUGAUGAG X CGAA ACUUGCUG	CAGCAAGT T TGAAGAA
3131	GUUCUCC CUGAUGAG X CGAA AACUUGCU	AGCAAGTT T GGAAGAAC
3147	GCAGGAAA CUGAUGAG X CGAA AUGUGGGG	CCCCACAT T TTCCTGCT
3148	CGCAGGAA CUGAUGAG X CGAA AAUGUGGG	CCCACATT T TTCCTGCG
3149	GCGCAGGA CUGAUGAG X CGAA AAAUGUGG	CCACATTT T TCCTGCGC
3150	CGCGCAGG CUGAUGAG X CGAA AAAAUGUG	CACATTTT T CCTGCGCG
3151	ACGCGCAG CUGAUGAG X CGAA AAAAAUGU	ACATTTT C CTGCGCGT
3160	UCAGAGAU CUGAUGAG X CGAA ACGCGCAG	CTGCGCGT C ATCTCTGA
3163	GUGUCAGA CUGAUGAG X CGAA AUGACGCG	CGCGTCAT C TGTGACAC
3165	CCGUGUCA CUGAUGAG X CGAA AGAUGACG	CGTCATCT C TGACACGG
3177	AGCAGAGG CUGAUGAG X CGAA AGGCCGUG	CACGGCCT C CCTCTGCT
3181	GAGUAGCA CUGAUGAG X CGAA AGGGAGGC	GCCTCCCT C TGCTACTC
3186	GGAUGGAG CUGAUGAG X CGAA AGCAGAGG	CCTCTGCT A CTCATCC
3189	UCAGGAUG CUGAUGAG X CGAA AGUAGCAG	CTGCTACT C CATCTGA
3193	GCUUUCAG CUGAUGAG X CGAA AUGGAGUA	TACTCCAT C CTGAAAGC
3219	CCCCCAGC CUGAUGAG X CGAA ACAUCCCU	AGGGATGT C GCTGGGGG
3248	GGAGGGCA CUGAUGAG X CGAA AGGGCCGG	CCGGCCCT C TGCCCTCC
3255	CGGCCUCG CUGAUGAG X CGAA AGGGCAGA	TCTGCCCT C CGAGGCCG
3288	UGAGCAGG CUGAUGAG X CGAA AUGCUUGG	CCAAGCAT T CCTGCTCA
3289	UUGAGCAG CUGAUGAG X CGAA AAUGCUUG	CAAGCAT T CTGCTCAA
3295	GUCAGCUU CUGAUGAG X CGAA AGCAGGAA	TTCCTGCT C AAGCTGAC
3305	ACGGUGUC CUGAUGAG X CGAA AGUCAGCU	AGCTGACT C GACACCGT
3316	ACGUAGGU CUGAUGAG X CGAA ACACGGUG	CACCGTGT C ACCTACGT
3321	GUGGCACG CUGAUGAG X CGAA AGGUGACA	TGTCACCT A CGTGCCAC
3331	GACCCAG CUGAUGAG X CGAA AGUGGCAC	GTGCCACT C CTGGGGTC
3339	UCCUGAGU CUGAUGAG X CGAA ACCCCAGG	CCTGGGGT C ACTCAGGA
3343	GCUGUCCU CUGAUGAG X CGAA AGUGACCC	GGGTCACT C AGGACAGC
3368	GAGCUUCC CUGAUGAG X CGAA ACUCAGCU	AGCTGAGT C GGAAGCTC
3376	GUCCCCGG CUGAUGAG X CGAA AGCUUCCG	CGGAAGCT C CCGGGGAC
3429	UGAAGUCU CUGAUGAG X CGAA AGGGCAGU	ACTGCCCT C AGACTTCA
3435	UGGUCUUG CUGAUGAG X CGAA AGUCUGAG	CTCAGACT T CAAGACCA
3436	AUGGUCUU CUGAUGAG X CGAA AAGUCUGA	TCAGACTT C AAGACCAT
3445	CAGUCCAG CUGAUGAG X CGAA AUGGUCUU	AAGACCAT C CTGGACTG
3503	CCCGGCGU CUGAUGAG X CGAA ACAGGGCU	AGCCCTGT C ACGCGGG
3514	GGGACGUA CUGAUGAG X CGAA AGCCCGGC	GCCGGGCT C TACGTCCC

Table 13

3516	CUGGGACG CUGAUGAG X CGAA AGAGCCCG	CGGGCTCT A CGTCCCAG
3520	CUCCCUUG CUGAUGAG X CGAA ACGUAGAG	CTCTACGT C CCAGGGAG
3568	AGGCCUCA CUGAUGAG X CGAA ACUCCCAG	CTGGGAGT C TGAGGCCT
3587	CUCGGCCA CUGAUGAG X CGAA ACACUCAC	GTGAGTGT T TGGCCGAG
3588	CCUCGGCC CUGAUGAG X CGAA AACACUCA	TGAGTGT T GGCCGAGG
3606	UUCAGCCG CUGAUGAG X CGAA ACAUGCAG	CTGCATGT C CGGCTGAA
3625	CUCAGCCG CUGAUGAG X CGAA ACACUCAG	CTGAGTGT C CGGCTGAG
3648	CUUGGCUG CUGAUGAG X CGAA ACACUCGC	GCGAGTGT C CAGCCAAG
3667	GUGUGCUG CUGAUGAG X CGAA ACACUCAG	CTGAGTGT C CAGCACAC
3683	GAAGUGAA CUGAUGAG X CGAA ACGGCAGG	CCTGCCGT C TTCCTTC
3685	GGGAAGUG CUGAUGAG X CGAA AGACGGCA	TGCCGTCT T CACTTCCC
3686	GGGAAGU CUGAUGAG X CGAA AAGACGGC	GCCGTCTT C ACTTCCCC
3690	CUGUGGG CUGAUGAG X CGAA AGUGAAGA	TCTTCACT T CCCACAG
3691	CCUGUGGG CUGAUGAG X CGAA AAGUGAAG	CTTCACTT C CCCACAGG
3708	GUGGAGCC CUGAUGAG X CGAA AGCGCCAG	CTGGCGCT C GGCTCCAC
3713	CUGGGGUG CUGAUGAG X CGAA AGCCGAGC	GCTCGGCT C CACCCAG
3730	GUGAGGAA CUGAUGAG X CGAA AGCUGGCC	GGCCAGCT T TTCTCAC
3731	GGUGAGGA CUGAUGAG X CGAA AAGCUGGC	GCCAGCTT T TCCTCACC
3732	UGGUGAGG CUGAUGAG X CGAA AAAGCUGG	CCAGCTTT T CCTCACCA
3733	CUGGUGAG CUGAUGAG X CGAA AAAAGCUG	CAGCTTTT C CTCACCAG
3736	CUCCUGGU CUGAUGAG X CGAA AGGAAAAG	CTTTTCCT C ACCAGGAG
3752	GGGAGUGG CUGAUGAG X CGAA AGCCGGGC	GCCCGGCT T CCACTCCC
3753	GGGGAGUG CUGAUGAG X CGAA AAGCCGGG	CCCGGCTT C CACTCCCC
3758	UAUGUGGG CUGAUGAG X CGAA AGUGGAAG	CTTCCACT C CCCACATA
3766	ACUAUUC CUGAUGAG X CGAA AUGUGGGG	CCCCACAT A GGAATAGT
3772	GGAUGGAC CUGAUGAG X CGAA AUUCCUAU	ATAGGAAT A GTCCATCC
3775	UGGGGAUG CUGAUGAG X CGAA ACUAUUC	GGAATAGT C CATCCCCA
3779	AAUCUGGG CUGAUGAG X CGAA AUGGACUA	TAGTCCAT C CCCAGATT
3787	CAAUGGCG CUGAUGAG X CGAA AUCUGGGG	CCCCAGAT T CGCCATTG
3788	ACAAUGGC CUGAUGAG X CGAA AAUCUGGG	CCCAGATT C GCCATTGT
3794	GGGUGAAC CUGAUGAG X CGAA AUGGCGAA	TTCCGCAT T GTTACCCC
3797	GAGGGGUG CUGAUGAG X CGAA ACAAUGGC	GCCATTGT T CACCCCTC
3798	CGAGGGGU CUGAUGAG X CGAA AACAAUGG	CCATTGTT C ACCCCTCG
3805	GGCAGGGC CUGAUGAG X CGAA AGGGGUGA	TCACCCCT C GGCCTGCC
3816	AGGCAAAG CUGAUGAG X CGAA AGGGCAGG	CCTGCCCT C CTTGCCT
3819	GGAAGGCA CUGAUGAG X CGAA AGGAGGGC	GCCCTCCT T TGCTTCC
3820	UGGAAGGC CUGAUGAG X CGAA AAGGAGGG	CCCTCCTT T GCCTTCCA
3825	GGGGGUGG CUGAUGAG X CGAA AGGCAAAG	CTTTGCCT T CCACCCCC
3826	UGGGGGUG CUGAUGAG X CGAA AAGGCAAA	TTTGCCTT C CACCCCCA
3839	UCCACCUG CUGAUGAG X CGAA AUGGUGGG	CCCACCAT C CAGGTGGA
3873	AAUUCCCA CUGAUGAG X CGAA AGCUCCCA	TGGGAGCT C TGGGAATT
3881	UCACUCCA CUGAUGAG X CGAA AUUCCCA	CTGGGAAT T TGGAGTGA
3882	GUCACUCC CUGAUGAG X CGAA AAUCCCA	TGGGAATT T GGAGTGAC
3907	CGCCUGUG CUGAUGAG X CGAA ACAGGGCA	TGCCCTGT A CACAGGCG
3940	CCCACAGG CUGAUGAG X CGAA ACCCCCAU	ATGGGGGT C CCTGTGGG
3950	CCCAAUUU CUGAUGAG X CGAA ACCCACAG	CTGTGGGT C AAATTGGG
3955	CUCCCCC CUGAUGAG X CGAA AUUGACC	GGTCAAAT T GGGGGGAG

Table 13

3977	CAGUAUUU CUGAUGAG X CGAA ACUCCCAC		GTGGGAGT A AAATACTG	
3982	AUAUUCAG CUGAUGAG X CGAA AUUUUACU		AGTAAAAT A CTGAATAT	
3989	AACUCAUA CUGAUGAG X CGAA AUUCAGUA		TACTGAAT A TATGAGTT	
3991	AAAACUCA CUGAUGAG X CGAA AUAUUCAG		CTGAATAT A TGAGTTTT	
3997	AACUGAAA CUGAUGAG X CGAA ACUCAUUA		ATATGAGT T TTTGAGTT	
3998	AAACUGAA CUGAUGAG X CGAA AACUCAUA		TATGAGTT T TTCAGTTT	
3999	AAAACUGA CUGAUGAG X CGAA AAACUCAU		ATGAGTTT T TCAGTTTT	
4000	CAAAACUG CUGAUGAG X CGAA AAAACUCA		TGAGTTTT T CAGTTTGT	
4001	UCAAACU CUGAUGAG X CGAA AAAACUC		GAGTTTTT C AGTTTTGA	
4005	UUUUUCAA CUGAUGAG X CGAA ACUGAAAA		TTTTCAGT T TTGAAAAA	
4006	UUUUUUA CUGAUGAG X CGAA AACUGAAA		TTTCAGTT T TGAAAAAA	
4007	UUUUUUUC CUGAUGAG X CGAA AAACUGAA		TTCAGTTT T GAAAAAAA	

Stem Length = 8 . Core Sequence = CUGAUGAG X CGAA (X = GCCGUUAGGC or other stem II sequence and length (greater than or equal to 2 base-pairs))

Seq1 = TERT (Homo sapiens telomerase reverse transcriptase (TERT) mRNA, 4015 bp); Nakamura *et al.*, Science 277 (5328), 955-959 (1997)

Table 14

Table 14: Human telomerase reverse transcriptase (TERT) NCH Ribozyme and Target Sequence

nt. Position	Ribozyme Sequence	Seq ID Nos	Substrate Sequence	Seq ID Nos
14	GCGCAGCA CUGAUGAG X CGAA IACGCAGC		GCTGCGTC C TGCTGCGC	
15	UGCGCAGC CUGAUGAG X CGAA IGACGCAG		CTGCGTCC T GCTGCGCA	
18	ACGUGCGC CUGAUGAG X CGAA ICAGGACG		CGTCCTGC T GCGCACGT	
23	UUCCACAG CUGAUGAG X CGAA ICGCAGCA		TGCTGCGC A CGTGGGAA	
34	GGGGCCAG CUGAUGAG X CGAA ICUUCCCA		TGGGAAGC C CTGGCCCC	
35	CGGGGCCA CUGAUGAG X CGAA IGCUCCCC		GGGAAGCC C TGGCCCCG	
36	CCGGGGCC CUGAUGAG X CGAA IGGCUUCC		GGAAGCCC T GGCCCCGG	
40	GUGGCCGG CUGAUGAG X CGAA ICCAGGGC		GCCCTGGC C CCGGCCAC	
41	GGUGGCCG CUGAUGAG X CGAA IGCCAGGG		CCCTGGCC C CGGCCACC	
42	GGUGGCC CUGAUGAG X CGAA IGGCCAGG		CCTGGCCC C GGCCACCC	
46	GCGGGGGU CUGAUGAG X CGAA ICCGGGGC		GCCCCGGC C ACCCCCGC	
47	CGCGGGGG CUGAUGAG X CGAA IGCCGGGG		CCCCGGCC A CCCCCGCG	
49	AUCGCGGG CUGAUGAG X CGAA IUGGCCGG		CCGGCCAC C CCGCGAT	
50	CAUCGCGG CUGAUGAG X CGAA IGUGGCCG		CGGCCACC C CCGCGATG	
51	GCAUCGCG CUGAUGAG X CGAA IGGUGGCC		GGCCACCC C CGCGATGC	
52	GGCAUCGC CUGAUGAG X CGAA IGGUGGCC		GCCACCCC C GCGATGCC	
60	GAGCGCGC CUGAUGAG X CGAA ICAUCGCG		CGCGATGC C GCGCGCTC	
67	CAGCGGGG CUGAUGAG X CGAA ICGCGCGG		CCGCGCGC T CCCCCTG	
69	GGCAGCGG CUGAUGAG X CGAA IAGCGCGC		GCGCGCTC C CCGCTGCC	
70	CGGAGCGG CUGAUGAG X CGAA IGAGCGCG		CGCGCTCC C CGCTGCCG	
71	UCGCGAGC CUGAUGAG X CGAA IGGAGCGC		GCGCTCCC C GCTGCCGA	
74	GGCUCGGC CUGAUGAG X CGAA ICGGGGAG		CTCCCGC T GCCGAGCC	
77	CACGGCUC CUGAUGAG X CGAA ICAGCGGG		CCCGCTGC C GAGCCGTG	
82	GAGCGCAC CUGAUGAG X CGAA ICUCGGCA		TGCCGAGC C GTGCGCTC	
89	CAGCAGGG CUGAUGAG X CGAA ICGCACGG		CCGTGCGC T CCCTGCTG	
91	CGCAGCAG CUGAUGAG X CGAA IAGCGCAC		GTGCGCTC C CTGCTGCG	
92	GCGCAGCA CUGAUGAG X CGAA IGAGCGCA		TGCGCTCC C TGCTGCGC	
93	UGCGCAGC CUGAUGAG X CGAA IGGAGCGC		GCGCTCCC T GCTGCGCA	
96	GGCUGCGC CUGAUGAG X CGAA ICAGGGAG		CTCCCTGC T GCGCAGCC	
101	GUAGUGGC CUGAUGAG X CGAA ICGCAGCA		TGCTGCGC A GCCACTAC	
104	GCGGUAGU CUGAUGAG X CGAA ICUGCGCA		TGCGCAGC C ACTACCGC	
105	CGCGUAG CUGAUGAG X CGAA IGCUGCGC		GCGCAGCC A CTACCGCG	
107	CUCGCGGU CUGAUGAG X CGAA IUGGCUGC		GCAGCCAC T ACGCGAG	
110	CACCUCGC CUGAUGAG X CGAA IUAGUGGC		GCCACTAC C GCGAGGTG	
120	CCAGCGGC CUGAUGAG X CGAA ICACCUCG		CGAGGTGC T GCCGCTGG	
123	UGGCCAGC CUGAUGAG X CGAA ICAGCACC		GGTGCTGC C GCTGGCCA	
126	ACGUGGCC CUGAUGAG X CGAA ICGGCAGC		GCTGCCGC T GGCCACGT	
130	ACGAACGU CUGAUGAG X CGAA ICCAGCGG		CCGCTGGC C ACGTTCGT	
131	CACGAACG CUGAUGAG X CGAA IGCCAGCG		CGCTGGCC A CGTTCGTG	
146	GGGCCCCA CUGAUGAG X CGAA ICGCCGCA		TGCGGCGC C TGGGGCCC	
147	GGGGCCCC CUGAUGAG X CGAA ICGCCGCG		GCGGCGCC T GGGGGCCC	
153	AGCCUUG CUGAUGAG X CGAA ICCCCAGG		CCTGGGGC C CCAGGGCT	
154	CAGCCUG CUGAUGAG X CGAA IGCCCCAG		CTGGGGCC C CAGGGCTG	

Table 14

155	CCAGCCCU CUGAUGAG X CGAA IGGCCCCA	TGGGGCCC C AGGGCTGG
156	GCCAGCCC CUGAUGAG X CGAA IGGGCCCC	GGGGCCCC A GGGCTGGC
161	CAGCCGCC CUGAUGAG X CGAA ICCUGGG	CCCAGGGC T GCGGCTG
168	GCUGCACC CUGAUGAG X CGAA ICCGCCAG	CTGGCGGC T GGTGCAGC
174	CCCCGCGC CUGAUGAG X CGAA ICACCAGC	GCTGGTGC A GCGCGGG
185	AGCCGCCG CUGAUGAG X CGAA IUCCCCGC	GCGGGAC C CGGCGCT
186	AAGCCGCC CUGAUGAG X CGAA IGUCCCCG	CGGGACC C GCGGCTT
193	GCGCGAA CUGAUGAG X CGAA ICCGCCG	CCGGCGGC T TTCCGCGC
197	CAGCGCGC CUGAUGAG X CGAA IAAAGCCG	CGGCTTTC C GCGCGCTG
204	GGGCCACC CUGAUGAG X CGAA ICGCGCGG	CCGCGCGC T GGTGGCCC
211	AGGCACUG CUGAUGAG X CGAA ICCACCAG	CTGGTGGC C CAGTGCCT
212	CAGGCACU CUGAUGAG X CGAA IGCCACCA	TGGTGGCC C AGTGCCTG
213	CCAGGCAC CUGAUGAG X CGAA IGGCCACC	GGTGGCCC A GTGCCTGG
218	GCACACCA CUGAUGAG X CGAA ICACUGGG	CCCAGTGC C TGGTGTGC
219	CGCACACC CUGAUGAG X CGAA IGCACUGG	CCAGTGCC T GGTGTGCG
231	CGUCCCAG CUGAUGAG X CGAA ICACGCAC	GTGCGTGC C CTGGACG
232	GCGUCCCA CUGAUGAG X CGAA IGCACGCA	TGCGTGCC C TGGACGC
233	UGCGUCCC CUGAUGAG X CGAA IGGCACGC	GCGTGCCC T GGGACGCA
241	GGCGGCCG CUGAUGAG X CGAA ICGUCCCA	TGGGACGC A CGGCCGCC
246	CGGGGGGC CUGAUGAG X CGAA ICCGUGCG	CGCACGGC C GCCCCCG
249	CGGCGGG CUGAUGAG X CGAA ICGGCCGU	ACGGCCGC C CCCCGCG
250	GCGGCGG CUGAUGAG X CGAA ICGGCCG	CGGCGCC C CCCCGCG
251	GGCGGCGG CUGAUGAG X CGAA IGGCGGCC	GGCGCCC C CCGCGCC
252	GGGCGGCG CUGAUGAG X CGAA IGGGCGGC	GCCGCCCC C CGCGCCC
253	GGGGCGGC CUGAUGAG X CGAA IGGGCGG	CCGCCCC C GCGCCCC
256	GAGGGGGC CUGAUGAG X CGAA ICGGGGG	CCCCCGC C GCCCCTC
259	AAGGAGG CUGAUGAG X CGAA ICGGCGG	CCGCGCC C CCTCTTC
260	GAAGGAGG CUGAUGAG X CGAA ICGGCGG	CCGCGCC C CCTCTTC
261	GGAAGGAG CUGAUGAG X CGAA IGGCGGCG	CGCGCCC C CTCCTTC
262	CGGAAGGA CUGAUGAG X CGAA IGGGCGGC	GCCGCCCC C TCCTTCG
263	GCGGAAGG CUGAUGAG X CGAA IGGGCGG	CCGCCCC T CTTCCGC
265	UGGCGGAA CUGAUGAG X CGAA IAGGGGGC	GCCCCCTC T TTCCGCA
266	CUGGCGGA CUGAUGAG X CGAA IGAGGGGG	CCCCCTC T TCCGCCAG
269	CACCUGGC CUGAUGAG X CGAA IAAGGAGG	CCTCTTC C GCCAGGTG
272	GGACACCU CUGAUGAG X CGAA ICGGAAGG	CCTTCGC C AGGTGTCC
273	AGGACACC CUGAUGAG X CGAA ICGGAAG	CTTCGCC A GGTGTCT
280	UUCAGGCA CUGAUGAG X CGAA IACACCU	CAGGTGTC C TGCCTGAA
281	CUUCAGGC CUGAUGAG X CGAA IGACACCU	AGGTGTCC T GCCTGAAG
284	CUCCUUCA CUGAUGAG X CGAA ICAGGACA	TGTCCTGC C TGAAGGAG
285	GCUCUUC CUGAUGAG X CGAA IGCAGGAC	GTCTGTC T GAAGGAGC
294	GGGCCACC CUGAUGAG X CGAA ICUCUUC	GAAGGAGC T GGTGGCCC
301	AGCACUC CUGAUGAG X CGAA ICCACCAG	CTGGTGGC C CGAGTGCT
302	CAGCACUC CUGAUGAG X CGAA IGCCACCA	TGGTGGCC C GAGTGCTG
309	GCCUCUGC CUGAUGAG X CGAA ICACUCGG	CCGAGTGC T GCAGAGGC
312	ACAGCCUC CUGAUGAG X CGAA ICAGCACU	AGTGCTGC A GAGGCTGT
318	GCUCGCAC CUGAUGAG X CGAA ICCUCUGC	GCAGAGGC T GTGCGAGC
345	CGAAGGCC CUGAUGAG X CGAA ICACGUUC	GAACGTGC T GGCCTTCG

Table 14

349	AAGCCGAA CUGAUGAG X CGAA ICCAGCAC		GTGCTGGC C TTCGGCTT	
350	GAAGCCGA CUGAUGAG X CGAA IGCCAGCA		TGCTGGCC T TCGGCTTC	
356	CAGCGCGA CUGAUGAG X CGAA ICCGAAGG		CCTTCGGC T TCGCGCTG	
363	CGUCCAGC CUGAUGAG X CGAA ICGGAAG		CTTCGCGC T GCTGGACG	
366	CCCCGUCC CUGAUGAG X CGAA ICAGCGCG		CGCGCTGC T GGACGGGG	
376	CCCCCGCG CUGAUGAG X CGAA ICCCCGUC		GACGGGGC C CGCGGGGG	
377	GCCCCCGC CUGAUGAG X CGAA IGCCCCGU		ACGGGGCC C GCGGGGGC	
386	CUCGGGGG CUGAUGAG X CGAA ICCCCCGC		GCGGGGGC C CCCCCGAG	
387	CCUCGGGG CUGAUGAG X CGAA IGCCCCCG		CGGGGGCC C CCCCCAGG	
388	GCCUCGGG CUGAUGAG X CGAA IGGCCCCC		GGGGGGCC C CCGAGGCC	
389	GGCCUCGG CUGAUGAG X CGAA IGGCCCCC		GGGGGGCC C CCGAGGCC	
390	AGGCCUCG CUGAUGAG X CGAA IGGGGCCC		GGGGGGCC C CGAGGCCT	
391	AAGGCCUC CUGAUGAG X CGAA IGGGGGCC		GGGGGGCC C GAGGCCTT	
397	GUGGUGAA CUGAUGAG X CGAA ICCUCGGG		CCCAGGCC C TTCACCAC	
398	GGUGGUGA CUGAUGAG X CGAA IGCCUCGG		CCGAGGCC T TCACCACC	
401	GCUGGUGG CUGAUGAG X CGAA IAAGCCCU		AGGCCTTC A CCACCAGC	
403	ACGCUGGU CUGAUGAG X CGAA IUGAAGGC		GCCTTCAC C ACCAGCGT	
404	CACGCUGG CUGAUGAG X CGAA IGUGAAGG		CCTTCACC A CCAGCGTG	
406	CGCACGCU CUGAUGAG X CGAA IUUGGUGAA		TTCACCAC C AGCGTGCG	
407	GCGCACGC CUGAUGAG X CGAA IGUGGUGA		TCACCACC A GCGTGCGC	
416	CAGGUAGC CUGAUGAG X CGAA ICGCACGC		GCGTGCGC A GCTACCTG	
419	GGGCAGGU CUGAUGAG X CGAA ICUGCGCA		TGCGCAGC T ACCTGCCC	
422	GUUGGGCA CUGAUGAG X CGAA IUAGCUGC		GCAGCTAC C TGCCCAAC	
423	UGUUGGGC CUGAUGAG X CGAA IGUAGCUG		CAGCTACC T GCCCAACA	
426	CCGUGUUG CUGAUGAG X CGAA ICAGGUAG		CTACCTGC C CAACACGG	
427	ACCGUGUU CUGAUGAG X CGAA IGCAGGUA		TACCTGCC C AACACGGT	
428	CACCGUGU CUGAUGAG X CGAA IGGCAGGU		ACCTGCCC A ACACGGTG	
431	GGUCACCG CUGAUGAG X CGAA IUUGGGCA		TGCCCAAC A CGGTGACC	
439	AGUGCGUC CUGAUGAG X CGAA IUCACCGU		ACGGTGAC C GACGACT	
445	CCCCGCAG CUGAUGAG X CGAA ICGUCGGU		ACCGACGC A CTGCGGGG	
447	UCCCCCGC CUGAUGAG X CGAA IUGCGUCG		CGACGCAC T GCGGGGGA	
471	GCAGCAGC CUGAUGAG X CGAA ICCCCCAC		GTGGGGGC T GCTGCTGC	
474	GGCGCAGC CUGAUGAG X CGAA ICAGCCCC		GGGGCTGC T GCTGCGCC	
477	CGCGGCGC CUGAUGAG X CGAA ICAGCAGC		GCTGCTGC T GCGCCGCG	
482	GCCACGC CUGAUGAG X CGAA ICGCAGCA		TGCTGCGC C GCGTGGGC	
501	GGUGAACC CUGAUGAG X CGAA ICAGUCG		CGACGTGC T GGTTCACC	
507	CCAGCAGG CUGAUGAG X CGAA IAACCAGC		GCTGGTTC A CCTGCTGG	
509	UGCCAGCA CUGAUGAG X CGAA IUGAACCA		TGGTTCAC C TGCTGGCA	
510	GUGCCAGC CUGAUGAG X CGAA IGUGAAC		GGTTCACC T GCTGGCAC	
513	AGCGUGCC CUGAUGAG X CGAA ICAGGUGA		TCACCTGC T GGCACGCT	
517	GCGCAGCG CUGAUGAG X CGAA ICCAGCAG		CTGCTGGC A CGTGCGC	
521	GAGCGCGC CUGAUGAG X CGAA ICGUGCCA		TGGCACGC T GCGCGCTC	
528	GCACAAAG CUGAUGAG X CGAA ICGCGCAG		CTGCGCGC T CTTGTGTC	
530	CAGCACAA CUGAUGAG X CGAA IAGCGCGC		GCGCGCTC T TTGTGCTG	
537	GAGCCACC CUGAUGAG X CGAA ICACAAAG		CTTTGTGC T GGTGGCTC	
544	CAGCUGGG CUGAUGAG X CGAA ICCACCAG		CTGGTGGC T CCCAGCTG	
546	CGCAGCUG CUGAUGAG X CGAA IAGCCACC		GGTGGCTC C CAGCTGCG	

Table 14

547	GCGCAGCU CUGAUGAG X CGAA IGAGCCAC		GTGGCTCC C AGCTGCGC	
548	GGCGCAGC CUGAUGAG X CGAA IGGAGCCA		TGGCTCCC A GCTGCGCC	
551	GUAGGCGC CUGAUGAG X CGAA ICUGGGAG		CTCCCAGC T GCGCTTAC	
556	ACCUGGUA CUGAUGAG X CGAA ICGCAGCU		AGCTGCGC C TACCAGGT	
557	CACCUGGU CUGAUGAG X CGAA ICGCAGC		GCTGCGCC T ACCAGGTG	
560	GCACACCU CUGAUGAG X CGAA IUAGGCGC		GCGCTTAC C AGGTGTGC	
561	CGCACACC CUGAUGAG X CGAA IGUAGGCG		CGCTTACC A GGTGTGCG	
573	ACAGCGGC CUGAUGAG X CGAA ICCCGCAC		GTGCGGGC C GCGCTGT	
576	GGUACAGC CUGAUGAG X CGAA ICGGCCCC		CGGGCCGC C GCTGTACC	
579	GCUGGUAC CUGAUGAG X CGAA ICGGCGGC		GCCGCGC T GTACCAGC	
584	GCCGAGCU CUGAUGAG X CGAA IUACAGCG		CGCTGTAC C AGCTCGGC	
585	CGCCGAGC CUGAUGAG X CGAA IGUACAGC		GCTGTACC A GCTCGGCG	
588	CAGCGCCG CUGAUGAG X CGAA ICUGGUAC		GTACCAGC T CGGCGCTG	
595	UGAGUGGC CUGAUGAG X CGAA ICGCCGAG		CTCGGCGC T GCCACTCA	
598	GCCUGAGU CUGAUGAG X CGAA ICAGCGCC		GGCGCTGC C ACTCAGGC	
599	GGCCUGAG CUGAUGAG X CGAA IGCAGCGC		GCGCTGCC A CTCAGGCC	
601	CGGGCCUG CUGAUGAG X CGAA IUGGCAGC		GCTGCCAC T CAGGCCCG	
603	GCCGGGCC CUGAUGAG X CGAA IAGUGGCA		TGCCACTC A GGCCCGGC	
607	GGGGGCCG CUGAUGAG X CGAA ICCUGAGU		ACTCAGGC C CGGCCCCC	
608	CGGGGGCC CUGAUGAG X CGAA IGCCUGAG		CTCAGGCC C GGCCCCCG	
612	GUGGCGGG CUGAUGAG X CGAA ICCGGGCC		GGCCCCGC C CCCGCCAC	
613	UGUGGCGG CUGAUGAG X CGAA IGCCGGGC		GCCCGGCC C CGCCACA	
614	GUGUGGCG CUGAUGAG X CGAA IGGCCGGG		CCCGGCC C CGCCACAC	
615	CGUGUGGC CUGAUGAG X CGAA IGGGCCGG		CCGGCCCC C GCCACAG	
618	UAGCGUGU CUGAUGAG X CGAA ICGGGGGC		GCCCCGC C ACACGCTA	
619	CUAGCGUG CUGAUGAG X CGAA ICGGGGGG		CCCCCGC A CACGCTAG	
621	CACUAGCG CUGAUGAG X CGAA IUGGCGGG		CCCCGCAC A CGCTAGTG	
625	GGUCCACU CUGAUGAG X CGAA ICGUGUGG		CCACACGC T AGTGGACC	
633	GCCUUCGG CUGAUGAG X CGAA IUCCACUA		TAGTGGAC C CCGAAGGC	
634	CGCCUUCG CUGAUGAG X CGAA IGUCCACU		AGTGGACC C CGAAGGCG	
635	ACGCCUUC CUGAUGAG X CGAA IGGUCCAC		GTGGACCC C GAAGGCGT	
645	CGCAUCCC CUGAUGAG X CGAA IACGCCUU		AAGGCGTC T GGGATGCG	
661	UGGUUCCA CUGAUGAG X CGAA ICCCGUUC		GAACGGGC C TGGAACCA	
662	AUGGUUCC CUGAUGAG X CGAA IGCCCGUU		AACGGGCC T GGAACCAT	
668	GACGCUAU CUGAUGAG X CGAA IUUCCAGG		CCTGGAAC C ATAGCGTC	
669	UGACGCUA CUGAUGAG X CGAA IGUCCAG		CTGGAACC A TAGCGTCA	
677	GGCCUCCC CUGAUGAG X CGAA IACGCUAU		ATAGCGTC A GGGAGGCC	
685	GGGACCCC CUGAUGAG X CGAA ICCUCCCU		AGGGAGGC C GGGGTCCC	
692	GCCCAGGG CUGAUGAG X CGAA IACCCCGG		CCGGGGTC C CCCTGGGC	
693	GGCCCAGG CUGAUGAG X CGAA IGACCCCG		CGGGGTCC C CCTGGGCC	
694	AGGCCCAG CUGAUGAG X CGAA IGGACCCC		GGGGTCCC C CTGGGCCT	
695	CAGGCCCA CUGAUGAG X CGAA IGGGACCC		GGGTCCCC C TGGCCTG	
696	GCAGGCCC CUGAUGAG X CGAA IGGGGACC		GGTCCCCC T GGGCCTGC	
701	GGCUGGCA CUGAUGAG X CGAA ICCCAGGG		CCCTGGGC C TGCCAGCC	
702	GGGUGGCG CUGAUGAG X CGAA IGCCCAGG		CCTGGGCC T GCCAGCCC	
705	CCGGGGCU CUGAUGAG X CGAA ICAGGCCC		GGGCTGCG C AGCCCCGG	
706	CCCGGGGC CUGAUGAG X CGAA IGCAGGCC		GGCTGCC A GCCCCGGG	

Table 14

709	GCACCCGG CUGAUGAG X CGAA ICUGGCAG	CTGCCAGC C CCGGTGTC
710	CGCACCCG CUGAUGAG X CGAA IGCUGGCA	TGCCAGCC C CCGGTGCG
711	UCGCACCC CUGAUGAG X CGAA IGGCUGGC	GCCAGCCC C GGGTGCGA
734	GCUGGCAC CUGAUGAG X CGAA ICCCCGC	GCGGGGGC A GTGCCAGC
739	CUUCGGCU CUGAUGAG X CGAA ICACUGCC	GGCAGTGC C AGCCGAAG
740	ACUUCGGC CUGAUGAG X CGAA IGCACUGC	GCAGTGCC A GCCGAAGT
743	CAGACUUC CUGAUGAG X CGAA ICUGGCAC	GTGCCAGC C GAAGTCTG
750	GCAACGGC CUGAUGAG X CGAA IACUUCGG	CCGAAGTC T GCCGTTGC
753	UGGGCAAC CUGAUGAG X CGAA ICAGACUU	AAGTCTGC C GTTGCCCA
759	GCCUCUUG CUGAUGAG X CGAA ICAACGGC	GCCGTTGC C CAAGAGGC
760	GGCCUCUU CUGAUGAG X CGAA IGCAACGG	CCGTGACC C AAGAGGCC
761	GGGCCUCU CUGAUGAG X CGAA IGGCAACG	CGTTGCCC A AGAGGCC
768	CACGCCUG CUGAUGAG X CGAA ICCUCUUG	CAAGAGGC C CAGGCTG
769	CCACGCCU CUGAUGAG X CGAA IGCCUCUU	AAGAGGCC C AGGCTGG
770	GCCACGCC CUGAUGAG X CGAA IGGCCUCU	AGAGGCC A GCGTGGC
781	UCAGGGGC CUGAUGAG X CGAA ICGCCACG	CGTGGCGC T GCCCTGA
784	GGCUCAGG CUGAUGAG X CGAA ICAGCGCC	GGCGTGC C CCTGAGCC
785	CGGCUCAG CUGAUGAG X CGAA IGCAGCGC	GCGTGCC C CTGAGCCG
786	CCGGCUCA CUGAUGAG X CGAA IGGCAGCG	CGCTGCCC C TGAGCCGG
787	UCCGGCUC CUGAUGAG X CGAA IGGGCAGC	GCTGCCCC T GAGCCGGA
792	UCCGCUCC CUGAUGAG X CGAA ICUCAGGG	CCCTGAGC C GGAGCGGA
804	GCCCAACG CUGAUGAG X CGAA ICGUCCGC	GCGGACGC C CGTTGGGC
805	UGCCCAAC CUGAUGAG X CGAA ICGGUCCG	CGGACGCC C GTTGGGCA
813	AGGACCCC CUGAUGAG X CGAA ICCCAACG	CGTTGGGC A GGGGTCTT
820	UGGGCCCA CUGAUGAG X CGAA IACCCUCG	CAGGGGTC C TGGGCCA
821	GUGGGCCC CUGAUGAG X CGAA IGACCCCU	AGGGGTCC T GGGCCAC
826	CCCGGGUG CUGAUGAG X CGAA ICCCAGGA	TCCTGGGC C CACCCGGG
827	GCCCGGGU CUGAUGAG X CGAA IGCCAGG	CCTGGGCC C ACCCGGGC
828	UGCCCGGG CUGAUGAG X CGAA IGGCCAG	CTGGGCCC A CCCGGGCA
830	CCUGCCCG CUGAUGAG X CGAA IUGGGCCC	GGGCCCAC C CGGGCAGG
831	UCCUGCCC CUGAUGAG X CGAA IGUGGGCC	GGCCCACC C GGGCAGGA
836	ACGCGUCC CUGAUGAG X CGAA ICCCGGGU	ACCCGGGC A GGACCGT
849	GGUCACUC CUGAUGAG X CGAA IUCCACGC	GCGTGAC C GAGTGACC
857	GAAACCAC CUGAUGAG X CGAA IUCACUCG	CGAGTGAC C GTGTTTC
866	CACCACAC CUGAUGAG X CGAA IAAACCAC	GTGTTTC T GTGTGTG
877	CUGGCAGG CUGAUGAG X CGAA IACACCAC	GTGTGTC A CCTGCCAG
879	GUCUGGCA CUGAUGAG X CGAA IUGACACC	GGTGTCAC C TGCCAGAC
880	GGUCUGGC CUGAUGAG X CGAA IGUGACAC	GTGTACAC T GCCAGACC
883	GCGGGUCU CUGAUGAG X CGAA ICAGGUGA	TCACCTGC C AGACCGC
884	GGCGGGUC CUGAUGAG X CGAA IGCAGGUG	CACCTGCC A GACCCGCC
888	CUUCGGCG CUGAUGAG X CGAA IUCUGGCA	TGCCAGAC C CGCCGAAG
889	UCUUCGGC CUGAUGAG X CGAA IGUCUGGC	GCCAGACC C GCCGAAGA
892	GCUUCUUC CUGAUGAG X CGAA ICGGGUCU	AGACCCGC C GAAGAAGC
901	AAAGAGGU CUGAUGAG X CGAA ICUUCUUC	GAAGAAGC C ACCTCTT
902	CAAAGAGG CUGAUGAG X CGAA IGCUCUUC	AAGAAGCC A CCTCTTG
904	UCCAAAGA CUGAUGAG X CGAA IUGGCUUC	GAAGCCAC C TCTTTGA
905	CUCCAAAG CUGAUGAG X CGAA IGUGGCUU	AAGCCACC T CTTTGAG

Table 14

907	CCCUCCAA CUGAUGAG X CGAA IAGGUGGC	GCCACCTC T TTGGAGGG
921	UGCCAGAG CUGAUGAG X CGAA ICGCACCC	GGGTGCGC T CTCTGGCA
923	CGUGCCAG CUGAUGAG X CGAA IAGCGCAC	GTGCGCTC T CTGGCACG
925	CGCGUGCC CUGAUGAG X CGAA IAGAGCGC	GCGCTCTC T GGCACGCG
929	GUGGCGCG CUGAUGAG X CGAA ICCAGAGA	TCTCTGGC A CGCGCCAC
935	GUGGGAGU CUGAUGAG X CGAA ICGCGUGC	GCACGCGC C ACTCCAC
936	GGUGGGAG CUGAUGAG X CGAA ICGCGUG	CACGCGCC A CTCCACC
938	UGGGUGGG CUGAUGAG X CGAA IUGGCGCG	CGCGCCAC T CCCACCA
940	GAUGGGUG CUGAUGAG X CGAA IAGUGGCG	CGCCACTC C CACCCATC
941	GGAUGGGU CUGAUGAG X CGAA IGAGUGGC	GCCACTCC C ACCCATCC
942	CGGAUGGG CUGAUGAG X CGAA IGGAGUGG	CCACTCCC A CCCATCCG
944	CACGGAUG CUGAUGAG X CGAA IUGGGAGU	ACTCCAC C CATCCGTG
945	CCACGGAU CUGAUGAG X CGAA IGUGGGAG	CTCCACC C ATCCGTGG
946	CCCACGGA CUGAUGAG X CGAA IGGUGGGA	TCCCACCC A TCCGTGGG
949	CGGCCAC CUGAUGAG X CGAA IAUGGGUG	CACCCATC C GTGGGCCG
956	GUGCUGGC CUGAUGAG X CGAA ICCACGG	CCGTGGGC C GCCAGCAC
959	GUGGUGCU CUGAUGAG X CGAA ICGGCCCA	TGGGCCGC C AGCACCAC
960	CGUGGUGC CUGAUGAG X CGAA ICGGCCC	GGGCCGCC A GCACCACG
963	CCGCGUGG CUGAUGAG X CGAA ICUGGCGG	CCGCCAGC A CCACGCGG
965	GCCCCGCU CUGAUGAG X CGAA IUGCUGGC	GCCAGCAC C ACGCGGGC
966	GGCCCCG CUGAUGAG X CGAA IGUGCUGG	CCAGCAC C CGCGGCC
974	GGAUGGGG CUGAUGAG X CGAA ICCCGCGU	ACGCGGGC C CCCATCC
975	UGGAUGGG CUGAUGAG X CGAA IGCCCGCG	CGCGGGCC C CCCATCCA
976	GUGGAUGG CUGAUGAG X CGAA IGGCCCGC	GCGGGCCC C CCATCCAC
977	UGUGGAUG CUGAUGAG X CGAA IGGGCCCG	CGGGCCCC C CATCCACA
978	AUGUGGAU CUGAUGAG X CGAA IGGGGCCC	GGGGCCCC C ATCCACAT
979	GAUGUGGA CUGAUGAG X CGAA IGGGGGCC	GGGGCCCC A TCCACATC
982	CGCGAUGU CUGAUGAG X CGAA IAUGGGGG	CCCCCATC C ACATCGCG
983	CCGCGAUG CUGAUGAG X CGAA IGAUGGGG	CCCCATCC A CATCGCGG
985	GGCCGCGA CUGAUGAG X CGAA IUGGAUGG	CCATCCAC A TCGCGGCC
993	GACGUGGU CUGAUGAG X CGAA ICCGCGAU	ATCGCGGC C ACCACGTC
994	GGACGUGG CUGAUGAG X CGAA ICGCGCGA	TCGCGGCC A CCACGTCC
996	AGGGACGU CUGAUGAG X CGAA IUGGCCGC	GCGGCCAC C ACGTCCCT
997	CAGGGACG CUGAUGAG X CGAA IGUGGCCG	CGGCCACC A CGTCCCTG
1002	UGUCCAG CUGAUGAG X CGAA IACGUGGU	ACCACGTC C CTGGGACA
1003	GUGUCCA CUGAUGAG X CGAA IGACGUGG	CCACGTCC C TGGGACAC
1004	CGUGUCCC CUGAUGAG X CGAA IGGACGUG	CACGTCCC T GGGACACG
1010	ACAAGGCG CUGAUGAG X CGAA IUCCAGG	CCTGGGAC A CGCCTTGT
1014	GGGGACAA CUGAUGAG X CGAA ICGUGUCC	GGACACGC C TTGTCCCC
1015	GGGGGACA CUGAUGAG X CGAA ICGUGUC	GACACGCC T TGTCCCC
1020	ACACCGGG CUGAUGAG X CGAA IACAAGGC	GCCTTGTC C CCCGGTGT
1021	UACACCGG CUGAUGAG X CGAA IGACAAGG	CCTTGTC C CCGGTGTA
1022	GUACACCG CUGAUGAG X CGAA IGGACAAG	CTTGTC C CGGTGTAC
1023	CGUACACC CUGAUGAG X CGAA IGGGACAA	TTGTCCCC C GGTGTACG
1033	UUGGUCUC CUGAUGAG X CGAA ICGUACAC	GTGTACGC C GAGACCAA
1039	AAGUGCUU CUGAUGAG X CGAA IUCUCGGC	GCCGAGAC C AAGCACTT
1040	GAAGUGCU CUGAUGAG X CGAA IGUCUCGG	CCGAGACC A AGCACTTC

Table 14

1044	AGAGGAAG CUGAUGAG X CGAA ICUUGGUC		GACCAAGC A CTTCTCT	
1046	GUAGAGGA CUGAUGAG X CGAA IUGCUUGG		CCAAGCAC T TCCTCTAC	
1049	GGAGUAGA CUGAUGAG X CGAA IAAGUGCU		AGCACTTC C TCTACTCC	
1050	AGGAGUAG CUGAUGAG X CGAA IGAAGUGC		GCACTTCC T CTACTCCT	
1052	UGAGGAGU CUGAUGAG X CGAA IAGGAAGU		ACTTCCTC T ACTCTCA	
1055	GCCUGAGG CUGAUGAG X CGAA IUAGAGGA		TCCTCTAC T CCTCAGGC	
1057	UCGCCUGA CUGAUGAG X CGAA IAGUAGAG		CTCTACTC C TCAGGCGA	
1058	GUCGCCUG CUGAUGAG X CGAA IGAGUAGA		TCTACTCC T CAGGCGAC	
1060	UUGUCGCC CUGAUGAG X CGAA IAGGAGUA		TACTCTC A GGCACAA	
1067	CUGCUCU CUGAUGAG X CGAA IUCGCCUG		CAGGCGAC A AGGAGCAG	
1074	GCCGCAGC CUGAUGAG X CGAA ICUCUUG		CAAGGAGC A GCTGCGGC	
1077	AGGGCCGC CUGAUGAG X CGAA ICUGCUC		GGAGCAGC T GCGGCCCT	
1083	GGAAGGAG CUGAUGAG X CGAA ICCGCAGC		GCTGCGGC C CTCCTCC	
1084	AGGAAGGA CUGAUGAG X CGAA IGCCGCAG		CTGCGGCC C TCCTCCT	
1085	UAGGAAGG CUGAUGAG X CGAA IGGCCGCA		TGCGGCC C CTTCTCTA	
1087	AGUAGGAA CUGAUGAG X CGAA IAGGGCCG		CGGCCCTC C TTCCTACT	
1088	GAGUAGGA CUGAUGAG X CGAA IGAGGGCC		GGCCCTCC T TCCTACTC	
1091	GCUGAGUA CUGAUGAG X CGAA IAAGGAGG		CCTCCTTC C TACTCAGC	
1092	AGCUGAGU CUGAUGAG X CGAA IGAAGGAG		CTCCTTC T ACTCAGCT	
1095	GAGAGCUG CUGAUGAG X CGAA IUAGGAAG		CTTCCTAC T CAGCTCTC	
1097	CAGAGAGC CUGAUGAG X CGAA IAGUAGGA		TCCTACTC A GCTCTCTG	
1100	CCUCAGAG CUGAUGAG X CGAA ICUGAGUA		TACTCAGC T CTCTGAGG	
1102	GGCCUCAG CUGAUGAG X CGAA IAGCUGAG		CTCAGCTC T CTGAGGCC	
1104	UGGGCCUC CUGAUGAG X CGAA IAGAGCUG		CAGCTCTC T GAGGCCCA	
1110	UCAGGCUG CUGAUGAG X CGAA ICCUCAGA		TCTGAGGC C CAGCCTGA	
1111	GUCAGGCU CUGAUGAG X CGAA IGCUCAG		CTGAGGCC C AGCCTGAC	
1112	AGUCAGGC CUGAUGAG X CGAA IGGCCUCA		TGAGGCC A GCCTGACT	
1115	GCCAGUCA CUGAUGAG X CGAA ICUGGGCC		GGCCAGC C TGA CTGGC	
1116	CGCCAGUC CUGAUGAG X CGAA IGCUGGGC		GCCAGCC T GACTGGCG	
1120	CGAGCGCC CUGAUGAG X CGAA IUCAGGCU		AGCCTGAC T GCGCTCG	
1126	AGCCUCCG CUGAUGAG X CGAA ICGCCAGU		ACTGGCGC T CGGAGGCT	
1134	UCUCCACG CUGAUGAG X CGAA ICCUCCGA		TCGGAGGC T CGTGAGA	
1144	AGAAAGAU CUGAUGAG X CGAA IUCUCCAC		GTGGAGAC C ATCTTTCT	
1145	CAGAAAGA CUGAUGAG X CGAA IGUCUCCA		TGGAGACC A TCTTTCTG	
1148	ACCCAGAA CUGAUGAG X CGAA IAUGGUCU		AGACCATC T TTCTGGGT	
1152	UGGAACCC CUGAUGAG X CGAA IAAAGAUG		CATCTTTC T GGGTTCCA	
1159	CAGGGCCU CUGAUGAG X CGAA IAACCCAG		CTGGGTTC C AGGCCCTG	
1160	CCAGGGCC CUGAUGAG X CGAA IGAACCCA		TGGGTTC A GGCCCTGG	
1164	GCAUCCAG CUGAUGAG X CGAA ICCUGGAA		TTCCAGGC C CTGGATGC	
1165	GGCAUCCA CUGAUGAG X CGAA IGCCUGGA		TCCAGGCC C TGGATGCC	
1166	UGGCAUCC CUGAUGAG X CGAA IGGCCUGG		CCAGGCC T GGATGCCA	
1173	GAGUCCU CUGAUGAG X CGAA ICAUCCAG		CTGGATGC C AGGGACTC	
1174	GGAGUCCC CUGAUGAG X CGAA IGCAUCCA		TGGATGCC A GGGACTCC	
1180	CUGCGGG CUGAUGAG X CGAA IUCCUGG		CCAGGGAC T CCGCAGG	
1182	ACCUGCG CUGAUGAG X CGAA IAGUCCU		AGGGACTC C CCGCAGGT	
1183	AACCUGCG CUGAUGAG X CGAA IGAGUCCC		GGGACTCC C CGCAGGTT	
1184	CAACCUGC CUGAUGAG X CGAA IGGAGUCC		GGACTCC C GCAGTTG	

Table 14

1187	GGGCAACC CUGAUGAG X CGAA ICGGGGAG	CTCCCCGC A GGTGCCCC
1194	GCAGGCGG CUGAUGAG X CGAA ICAACCUG	CAGGTTGC C CCGCCTGC
1195	GGCAGGCG CUGAUGAG X CGAA IGCAACCU	AGGTTGCC C CGCCTGCC
1196	GGGCAGGC CUGAUGAG X CGAA IGGCAACC	GGTTGCCC C GCCTGCCC
1199	CUGGGGCA CUGAUGAG X CGAA ICGGGGCA	TGCCCCGC C TGCCCCAG
1200	GCUGGGGC CUGAUGAG X CGAA ICGGGGCG	GCCCCGCC T GCCCCAGC
1203	AGCGCUGG CUGAUGAG X CGAA ICAGGCGG	CCGCCTGC C CCAGCGCT
1204	UAGCGCUG CUGAUGAG X CGAA IGCAGGCG	CGCCTGCC C CAGCGCTA
1205	GUAGCGCU CUGAUGAG X CGAA IGGCAGGC	GCCTGCCC C AGCGCTAC
1206	AGUAGCGC CUGAUGAG X CGAA IGGCAGG	CCTGCCCC A GCGCTACT
1211	UUGCCAGU CUGAUGAG X CGAA ICGCUGGG	CCCAGCGC T ACTGGCAA
1214	CAUUUGCC CUGAUGAG X CGAA IUAGCGCU	AGCGCTAC T GGCAAATG
1218	GCCGCAUU CUGAUGAG X CGAA ICCAGUAG	CTACTGGC A AATGCGGC
1227	GAAACAGG CUGAUGAG X CGAA ICCGCAUU	AATGCGGC C CCTGTTTC
1228	AGAAACAG CUGAUGAG X CGAA IGCCGCAU	ATGCGGCC C CTGTTTCT
1229	CAGAAACA CUGAUGAG X CGAA IGGCCGCA	TGCGGCCC C TGTTCCTG
1230	CCAGAAAC CUGAUGAG X CGAA IGGGCCGC	GCGGCCCC T GTTTCCTG
1236	GCAGCUCC CUGAUGAG X CGAA IAAACAGG	CCTGTTTC T GGAGCTGC
1242	UCCCAAGC CUGAUGAG X CGAA ICUCCAGA	TCTGGAGC T GCTTGGGA
1245	GGUUCCCA CUGAUGAG X CGAA ICAGCUCC	GGAGCTGC T TGGGAACC
1253	CUGCGCGU CUGAUGAG X CGAA IUUCCCAA	TTGGGAAC C ACGCGCAG
1254	ACUGCGCG CUGAUGAG X CGAA IGUUCCCA	TGGGAACC A CGCGCAGT
1260	AGGGGCAC CUGAUGAG X CGAA ICGCGUGG	CCACGCGC A GTGCCCT
1265	CCCGUAGG CUGAUGAG X CGAA ICACUGCG	CGCAGTGC C CCTACGGG
1266	CCCCGUAG CUGAUGAG X CGAA IGCACUGC	GCAGTGCC C CTACGGGG
1267	ACCCCGUA CUGAUGAG X CGAA IGGCACUG	CAGTGCCC C TACGGGGT
1268	CACCCCGU CUGAUGAG X CGAA IGGGCACU	AGTGCCCC T ACGGGGTG
1278	UCUUGAGG CUGAUGAG X CGAA ICACCCCG	CGGGGTGC T CCTCAAGA
1280	CGUCUUGA CUGAUGAG X CGAA IAGCACCC	GGGTGCTC C TCAAGACG
1281	GCGUCUUG CUGAUGAG X CGAA IGAGCACC	GGTGCTCC T CAAGACGC
1283	GUGCGUCU CUGAUGAG X CGAA IAGGAGCA	TGCTCCTC A AGACGCAC
1290	GCGGGCAG CUGAUGAG X CGAA ICGUCUUG	CAAGACGC A CTGCCCCG
1292	CAGCGGGC CUGAUGAG X CGAA IUGCGUCU	AGACGCAC T GCCCGCTG
1295	UCGCAGCG CUGAUGAG X CGAA ICAGUGCG	CGCACTGC C CGCTGCGA
1296	CUCGCAGC CUGAUGAG X CGAA IGCAGUGC	GCACTGCC C GCTGCGAG
1299	CAGCUCGC CUGAUGAG X CGAA ICGGGCAG	CTGCCCCG T GCGAGCTG
1306	GUGACCGC CUGAUGAG X CGAA ICUCGCAG	CTGCGAGC T GCGGTCAC
1313	UGCUGGGG CUGAUGAG X CGAA IACCGCAG	CTGCGGTC A CCCCAGCA
1315	GCUGCUGG CUGAUGAG X CGAA IUGACCGC	GCGGTCAC C CCAGCAGC
1316	GGCUGCUG CUGAUGAG X CGAA IGUGACCG	CGGTACCC C CAGCAGCC
1317	CGGCUGCU CUGAUGAG X CGAA IGGUGACC	GGTCACCC C AGCAGCCG
1318	CCGGCUGC CUGAUGAG X CGAA IGGGUGAC	GTCACCCC A GCAGCCGG
1321	ACACCCGC CUGAUGAG X CGAA ICUGGGGU	ACCCAGC A GCCGGTGT
1324	CAGACACC CUGAUGAG X CGAA ICUGCUGG	CCAGCAGC C GGTGTCTG
1331	CCGGGCAC CUGAUGAG X CGAA IACACCGG	CCGGTGTC T GTGCCCGG
1336	UUCUCCCG CUGAUGAG X CGAA ICACAGAC	GTCTGTGC C CGGGAGAA
1337	CUUCUCCC CUGAUGAG X CGAA IGCACAGA	TCTGTGCC C GGGAGAAG

Table 14

1347	AGCCCUUG CUGAUGAG X CGAA ICUUCUCC		GGAGAAGC C CCAGGGCT	
1348	GAGCCUUG CUGAUGAG X CGAA IGCUCUC		GAGAAGCC C CAGGGCTC	
1349	AGAGCCCU CUGAUGAG X CGAA IGGCUUCU		AGAAGCCC C AGGGCTCT	
1350	CAGAGCCC CUGAUGAG X CGAA IGGCUUC		GAAGCCCC A GGGCTCTG	
1355	CGCCACAG CUGAUGAG X CGAA ICCUGGG		CCCAGGGC T CTGTGGCG	
1357	GCCGCCAC CUGAUGAG X CGAA IAGCCUG		CAGGGCTC T GTGGCGGC	
1366	UCCUCGGG CUGAUGAG X CGAA ICCGCCAC		GTGGCGGC C CCCGAGGA	
1367	CUCCUCGG CUGAUGAG X CGAA IGCCGCCA		TGGCGGCC C CCCGAGGAG	
1368	CCUCCUCG CUGAUGAG X CGAA IGGCCGCC		GGCGGCC C CGAGGAGG	
1369	UCCUCCUC CUGAUGAG X CGAA IGGCCGCG		GCGGCC C GAGGAGGA	
1382	GGGUCUG CUGAUGAG X CGAA IUCCUCU		AGGAGGAC A CAGACCCC	
1384	CGGGGUC CUGAUGAG X CGAA IUUGCCUC		GAGGACAC A GACCCCG	
1388	GCGACGG CUGAUGAG X CGAA IUCUGUGU		ACACAGAC C CCCGTCGC	
1389	GGCGACGG CUGAUGAG X CGAA IGUCUGUG		CACAGACC C CCGTCGCC	
1390	AGCGACG CUGAUGAG X CGAA IGGUCUGU		ACAGACC C CGTCGCT	
1391	CAGCGAC CUGAUGAG X CGAA IGGUCUG		CAGACC C GTCGCTG	
1397	CUGCACCA CUGAUGAG X CGAA ICGACGGG		CCCGTCGC C TGGTGACG	
1398	GCUGCACC CUGAUGAG X CGAA ICGACGG		CCGTCGCC T GGTGCAGC	
1404	GGAGCAGC CUGAUGAG X CGAA ICACCAGG		CCTGGTGC A GCTGCTCC	
1407	GGCGAGC CUGAUGAG X CGAA ICUGCACC		GGTGACG T GCTCCGCC	
1410	GCUGGCG CUGAUGAG X CGAA ICAGCUGC		GCAGCTGC T CCGCCAGC	
1412	GUGCUGGC CUGAUGAG X CGAA IAGCAGCU		AGCTGCTC C GCCAGCAC	
1415	GCUGUGCU CUGAUGAG X CGAA ICGGAGCA		TGCTCCGC C AGCACAGC	
1416	UGCUGUC CUGAUGAG X CGAA ICGGAGC		GCTCCGCC A GCACAGCA	
1419	GGCUGUC CUGAUGAG X CGAA ICUGGCGG		CCGCCAGC A CAGCAGCC	
1421	GGGUCUC CUGAUGAG X CGAA IUGCUGGC		GCCAGCAC A GCAGCCCC	
1424	CCAGGGC CUGAUGAG X CGAA ICUGUGCU		AGCACAGC A GCCCTGG	
1427	CUGCCAG CUGAUGAG X CGAA ICUGCUGU		ACAGCAGC C CCTGGCAG	
1428	CCUGCCAG CUGAUGAG X CGAA IGCUGCUG		CAGCAGCC C CTGGCAGG	
1429	ACCUGCCA CUGAUGAG X CGAA IGGCUGCU		AGCAGCCC C TGGCAGGT	
1430	CACCUGCC CUGAUGAG X CGAA IGGCUGC		GCAGCCCC T GGCAGGTG	
1434	CGUACACC CUGAUGAG X CGAA ICCAGGGG		CCCCTGGC A GGTGTACG	
1445	CCGCACGA CUGAUGAG X CGAA ICCGUACA		TGTACGGC T TCGTCGG	
1456	CGCAGGCA CUGAUGAG X CGAA ICCGCAC		GTGCGGGC C TGCCTGCG	
1457	GCGCAGC CUGAUGAG X CGAA IGCCGCA		TGCGGGC T GCCTGCGC	
1460	CCGGCGCA CUGAUGAG X CGAA ICAGGCC		GGCCTGC C TGCCTGG	
1461	GCGGCGC CUGAUGAG X CGAA IGCAGGC		GGCCTGCC T GCGCCGGC	
1466	CACCAGCC CUGAUGAG X CGAA ICGCAGGC		GCTGCGC C GGCTGGTG	
1470	GGGACCC CUGAUGAG X CGAA ICCGGCGC		GCGCCGGC T GGTGCCCC	
1476	GGCCUGGG CUGAUGAG X CGAA ICACCAGC		GCTGGTGC C CCCAGGCC	
1477	AGGCCUG CUGAUGAG X CGAA IGCACCAG		CTGGTGCC C CCAGGCCT	
1478	GAGGCCUG CUGAUGAG X CGAA IGGCACCA		TGGTGCCC C CAGGCCTC	
1479	AGAGGCCU CUGAUGAG X CGAA IGGCACCC		GGTGCCCC C AGGCCTCT	
1480	CAGAGGCC CUGAUGAG X CGAA IGGGCAC		GTGCCCCC A GGCTCTG	
1484	GCCCCAGA CUGAUGAG X CGAA ICCUGGGG		CCCCAGGC C TCTGGGGC	
1485	AGCCCCAG CUGAUGAG X CGAA IGCCUGGG		CCCAGGCC T CTGGGGCT	
1487	GGAGCCCC CUGAUGAG X CGAA IAGGCCUG		CAGGCCTC T GGGGCTCC	

Table 14

1493	GUGCCUGG CUGAUGAG X CGAA ICCCCAGA		TCTGGGGC T CCAGGCAC	
1495	UUGUGCCU CUGAUGAG X CGAA IAGCCCCA		TGGGGCTC C AGGCACAA	
1496	GUUGUGCC CUGAUGAG X CGAA IGAGCCCC		GGGGCTCC A GGCACAA	
1500	GUUCGUUG CUGAUGAG X CGAA ICCUGGAG		CTCCAGGC A CAACGAAC	
1502	GCGUUCGU CUGAUGAG X CGAA IUGCCUGG		CCAGGCAC A ACGAACGC	
1511	GAGGAAGC CUGAUGAG X CGAA ICGUUCGU		ACGAACGC C GCTTCCTC	
1514	CCUGAGGA CUGAUGAG X CGAA ICGGCGUU		AACGCCGC T TCCTCAGG	
1517	GUUCCUGA CUGAUGAG X CGAA IAAGCGGC		GCCGCTTC C TCAGGAAC	
1518	UGUCCUG CUGAUGAG X CGAA IGAAGCGG		CCGCTTCC T CAGGAACA	
1520	GGUGUCC CUGAUGAG X CGAA IAGGAAGC		GCTTCCTC A GGAACACC	
1526	CUUCUUGG CUGAUGAG X CGAA IUUCCUGA		TCAGGAAC A CCAAGAAG	
1528	AACUUCUU CUGAUGAG X CGAA IUGUCCU		AGGAACAC C AAGAAGTT	
1529	GAACUUCU CUGAUGAG X CGAA IGUGUCC		GGAACACC A AGAAGTTC	
1538	CAGGGAGA CUGAUGAG X CGAA IAACUUCU		AGAAGTTC A TCTCCCTG	
1541	CCCCAGGG CUGAUGAG X CGAA IAUGAACU		AGTTCATC T CCCTGGGG	
1543	UUCCCCAG CUGAUGAG X CGAA IAGAUGAA		TTCATCTC C CTGGGGAA	
1544	CUUCCCCA CUGAUGAG X CGAA IGAGAUGA		TCATCTCC C TGGGGAAG	
1545	GCUUCCCC CUGAUGAG X CGAA IGGAGAUG		CATCTCCC T GGGGAAGC	
1554	GCUUGGCA CUGAUGAG X CGAA ICUUCCCC		GGGGAAGC A TGCCAAGC	
1558	GAGAGCUU CUGAUGAG X CGAA ICAUGCUU		AAGCATGC C AAGCTCTC	
1559	CGAGAGCU CUGAUGAG X CGAA IGCAUGCU		AGCATGCC A AGCTCTCG	
1563	GCAGCGAG CUGAUGAG X CGAA ICUUGGCA		TGCCAAGC T CTCGCTGC	
1565	CUGCAGCG CUGAUGAG X CGAA IAGCUUGG		CCAAGCTC T CGCTGCAG	
1569	GUCCUGC CUGAUGAG X CGAA ICGAGAGC		GCTCTCGC T GCAGGAGC	
1572	UCAGCUCC CUGAUGAG X CGAA ICAGCGAG		CTCGCTGC A GGAGCTGA	
1578	UCCACGUC CUGAUGAG X CGAA ICUCCUGC		GCAGGAGC T GACGTGGA	
1604	CCAAGCGC CUGAUGAG X CGAA IUCCCGCA		TGCGGGAC T GCGCTGG	
1609	CGCAGCCA CUGAUGAG X CGAA ICGCAGUC		GA CTGCGC T TGGCTGCG	
1614	UCCUGCGC CUGAUGAG X CGAA ICCAAGCG		CGCTTGGC T GCGCAGGA	
1619	UGGGCUCC CUGAUGAG X CGAA ICGCAGCC		GGCTGCGC A GGAGCCCA	
1625	AACCCUG CUGAUGAG X CGAA ICUCCUGC		GCAGGAGC C CAGGGTTT	
1626	CAACCCU CUGAUGAG X CGAA IGCUCUG		CAGGAGCC C AGGGTTG	
1627	CCAACCC CUGAUGAG X CGAA IGGUCCU		AGGAGCCC A GGGTTGG	
1637	CGGAACAC CUGAUGAG X CGAA ICCAACCC		GGGTGGC T GTGTTCCG	
1644	CUGCGGCC CUGAUGAG X CGAA IAACACAG		CTGTGTTC C GGCCGAG	
1648	UGCUCUGC CUGAUGAG X CGAA ICCGGAAC		GTTCCGGC C GCAGAGCA	
1651	CGGUGCUC CUGAUGAG X CGAA ICGGCCGG		CCGGCCGC A GAGACCG	
1656	GCAGACGG CUGAUGAG X CGAA ICUCUGCG		CGCAGAGC A CCGTCTGC	
1658	ACGACAGC CUGAUGAG X CGAA IUGCUCUG		CAGAGCAC C GTCTGCGT	
1662	CCUCACGC CUGAUGAG X CGAA IACGGUGC		GCACCGTC T GCGTGAGG	
1676	CUUGGCCA CUGAUGAG X CGAA IAUCCU		AGGAGATC C TGGCCAAG	
1677	ACUUGGCC CUGAUGAG X CGAA IGAUCCU		GGAGATCC T GGCCAAGT	
1681	AGGAACUU CUGAUGAG X CGAA ICCAGGAU		ATCCTGGC C AAGTTCCT	
1682	CAGGAACU CUGAUGAG X CGAA IGCCAGGA		TCCTGGCC A AGTTCCTG	
1688	CCAGUGCA CUGAUGAG X CGAA IAACUUGG		CCAAGTTC C TGCACTGG	
1689	GCCAGUGC CUGAUGAG X CGAA IGAACUUG		CAAGTTC T GCACTGGC	
1692	UCAGCCAG CUGAUGAG X CGAA ICAGGAAC		GTTCTGCG A CTGGCTGA	

Table 14

1694	CAUCAGCC CUGAUGAG X CGAA IUGCAGGA	TCCTGCAC T GGCTGATG
1698	CACUCAUC CUGAUGAG X CGAA ICCAGUGC	GCACTGGC T GATGAGTG
1722	ACCUGAGC CUGAUGAG X CGAA ICUCGACG	CGTCGAGC T GCTCAGGT
1725	AAGACCUG CUGAUGAG X CGAA ICAGCUCG	CGAGCTGC T CAGGTCTT
1727	GAAAGACC CUGAUGAG X CGAA IAGCAGCU	AGCTGCTC A GGTCTTTC
1732	UAAAAGAA CUGAUGAG X CGAA IACCUGAG	CTCAGGTC T TTCTTTTA
1736	GACAUAAA CUGAUGAG X CGAA IAAAGACC	GGTCTTTC T TTTATGTC
1745	GGUCUCCG CUGAUGAG X CGAA IACAUAAA	TTTATGTC A CGGAGACC
1753	UGAAACGU CUGAUGAG X CGAA IUUCUCCGU	ACGGAGAC C ACGTTTCA
1754	UUGAAACG CUGAUGAG X CGAA IGUCUCCG	CGGAGACC A CGTTTCAA
1761	UGUUCUUU CUGAUGAG X CGAA IAAACGUG	CACGTTTC A AAAGAACA
1769	AAAGAGCC CUGAUGAG X CGAA IUUCUUUU	AAAAGAAC A GGCTCTTT
1773	AGAAAAAG CUGAUGAG X CGAA ICCUGUUC	GAACAGGC T CTTTTTCT
1775	GUAGAAAA CUGAUGAG X CGAA IAGCCUGU	ACAGGCTC T TTTTCTAC
1781	CUUCCGGU CUGAUGAG X CGAA IAAAAAGA	TCTTTTTC T ACCGGAAG
1784	ACUCUUC CUGAUGAG X CGAA IUAGAAAA	TTTTCTAC C GGAAGAGT
1796	CUUGCUC CUGAUGAG X CGAA IACACUCU	AGAGTGTC T GGAGCAAG
1802	UUGCAACU CUGAUGAG X CGAA ICUCCAGA	TCTGGAGC A AGTTGCAA
1809	CAAUGCUU CUGAUGAG X CGAA ICAACUUG	CAAGTTGC A AAGCATTG
1814	GAUUCCAA CUGAUGAG X CGAA ICUUUGCA	TGCAAAGC A TTGAATC
1823	GUGCUGUC CUGAUGAG X CGAA IAUUCCAA	TTGAATC A GACAGCAC
1827	UCAAGUGC CUGAUGAG X CGAA IUCUGAUU	AATCAGAC A GCACTTGA
1830	UCUUCAAG CUGAUGAG X CGAA ICUGUCUG	CAGACAGC A CTTGAAGA
1832	CCUCUUCA CUGAUGAG X CGAA IUGCUGUC	GACAGCAC T TGAAGAGG
1845	CCCGCAGC CUGAUGAG X CGAA ICACCCUC	GAGGGTGC A GCTGCGGG
1848	GCUCCCGC CUGAUGAG X CGAA ICUGCACC	GGTGCAGC T GCGGGAGC
1857	CUUCCGAC CUGAUGAG X CGAA ICUCCCGC	GCGGGAGC T GTCGGAAG
1867	CUGACCUC CUGAUGAG X CGAA ICUUCGCA	TCGGAAGC A GAGGTCAG
1874	AUGCUGCC CUGAUGAG X CGAA IACCUCUG	CAGAGGTC A GGCAGCAT
1878	CCCGAUGC CUGAUGAG X CGAA ICCUGACC	GGTCAGGC A GCATCGGG
1881	CUUCCCGA CUGAUGAG X CGAA ICUGCCUG	CAGGCAGC A TCGGGAAG
1891	GCGGGCCU CUGAUGAG X CGAA ICUCCCCG	CGGGAAGC C AGGCCCGC
1892	GGCGGGCC CUGAUGAG X CGAA IGCUCCCC	GGGAAGCC A GGCCCCGC
1896	GCAGGGCG CUGAUGAG X CGAA ICCUGGCU	AGCCAGGC C CGCCCTGC
1897	AGCAGGGC CUGAUGAG X CGAA IGCCUGGC	GCCAGGCC C GCCCTGCT
1900	GUCAGCAG CUGAUGAG X CGAA ICGGGCCU	AGGCCCGC C CTGCTGAC
1901	CGUCAGCA CUGAUGAG X CGAA IGCGGGCC	GGCCCGCC C TGCTGACG
1902	ACGUCAGC CUGAUGAG X CGAA IGGCGGGC	GCCCGCCC T GCTGACGT
1905	UGGACGUC CUGAUGAG X CGAA ICAGGGCG	CGCCCTGC T GACGTCCA
1912	CGGAGUCU CUGAUGAG X CGAA IACGUCAG	CTGACGTC C AGACTCCG
1913	GCGGAGUC CUGAUGAG X CGAA IGACGUCA	TGACGTCC A GACTCCGC
1917	UGAAGCGG CUGAUGAG X CGAA IUCUGGAC	GTCCAGAC T CCGTTTCA
1919	GAUGAAGC CUGAUGAG X CGAA IAGUCUGG	CCAGACTC C GCTTCATC
1922	GGGAUGA CUGAUGAG X CGAA ICGGAGUC	GACTCCGC T TCATCCCC
1925	CUUGGGGA CUGAUGAG X CGAA IAAGCGGA	TCCGCTTC A TCCCCAAG
1928	AGGCUUGG CUGAUGAG X CGAA IAUGAAGC	GCTTCATC C CCAAGCCT
1929	CAGGCUUG CUGAUGAG X CGAA IGAUGAAG	CTTCATCC C CAAGCCTG

Table 14

1930	UCAGGCUU CUGAUGAG X CGAA IGGAUGAA	TTCATCCC C AAGCCTGA
1931	GUCAGGCU CUGAUGAG X CGAA IGGGAUGA	TCATCCCC A AGCCTGAC
1935	GCCCGUCA CUGAUGAG X CGAA ICUUGGGG	CCCCAAGC C TGACGGGC
1936	AGCCCGUC CUGAUGAG X CGAA IGCUUGGG	CCCAAGCC T GACGGGCT
1944	UCGGCCGC CUGAUGAG X CGAA ICCCGUCA	TGACGGGC T GCGGCCGA
1950	UCACAAUC CUGAUGAG X CGAA ICCGCAGC	GCTGCGGC C GATTGTGA
1961	GUAGUCCA CUGAUGAG X CGAA IUUCACAA	TTGTGAAC A TGGACTAC
1967	CACGACGU CUGAUGAG X CGAA IUCCAUGU	ACATGGAC T ACGTCGTG
1981	AACGUUCU CUGAUGAG X CGAA ICUCCAC	GTGGGAGC C AGAACGTT
1982	GAACGUUC CUGAUGAG X CGAA IGCUCCCA	TGGGAGCC A GAACGTTT
1991	UUCUCUGC CUGAUGAG X CGAA IAACGUUC	GAACGTTT C GCAGAGAA
1994	CUUUUCUC CUGAUGAG X CGAA ICGGAACG	CGTTCCGC A GAGAAAAG
2008	AGACGCUC CUGAUGAG X CGAA ICCUCUU	AAGAGGGC C GAGCGTCT
2016	UCGAGGUG CUGAUGAG X CGAA IACGCUCG	CGAGCGTC T CACCTCGA
2018	CCUCGAGG CUGAUGAG X CGAA IAGACGCU	AGCGTCTC A CCTCGAGG
2020	ACCCUCGA CUGAUGAG X CGAA IUGAGACG	CGTCTCAC C TCGAGGGT
2021	CACCCUCG CUGAUGAG X CGAA IGUGAGAC	GTCTCACC T CGAGGGTG
2035	CUGAACAG CUGAUGAG X CGAA ICCUUCAC	GTGAAGGC A CTGTTTCA
2037	CGCUGAAC CUGAUGAG X CGAA IUGCCUUC	GAAGGCAC T GTTCAGCG
2042	GAGCACGC CUGAUGAG X CGAA IAACAGUG	CACTGTTC A GCGTGCTC
2049	CGUAGUUG CUGAUGAG X CGAA ICACGCUG	CAGCGTGC T CAACTACG
2051	CUCGUAGU CUGAUGAG X CGAA IAGCACGC	GCGTGCTC A ACTACGAG
2054	CCGCUCGU CUGAUGAG X CGAA IUUGAGCA	TGCTCAAC T ACGAGCGG
2072	GAGGCCGG CUGAUGAG X CGAA ICGCCGCG	CGCGGCGC C CCGCCTC
2073	GGAGGCCG CUGAUGAG X CGAA ICGCCGCG	GCGGCGCC C CGGCCTC
2074	AGGAGGCC CUGAUGAG X CGAA IGGCGCCG	CGGCGCCC C GGCCTCCT
2078	GCCCAGGA CUGAUGAG X CGAA ICCGGGGC	GCCCCGGC C TCCTGGGC
2079	CGCCAGG CUGAUGAG X CGAA IGCCGGGG	CCCCGGCC T CCTGGGCG
2081	GGCGCCA CUGAUGAG X CGAA IAGGCCGG	CCGGCCTC C TGGGCGCC
2082	AGGCGCCC CUGAUGAG X CGAA IGAGGCCG	CGGCCTCC T GGGCGCCT
2089	AGCACAGA CUGAUGAG X CGAA ICGCCAG	CTGGGCGC C TCTGTGCT
2090	CAGCACAG CUGAUGAG X CGAA ICGGCCCA	TGGGCGCC T CTGTGCTG
2092	CCCAGCAC CUGAUGAG X CGAA IAGGCGCC	GGCGCCTC T GTGCTGGG
2097	CCAGGCCC CUGAUGAG X CGAA ICACAGAG	CTCTGTGC T GGGCCTGG
2102	AUCGUCCA CUGAUGAG X CGAA ICCAGCA	TGCTGGGC C TGGACGAT
2103	UAUCGUCC CUGAUGAG X CGAA IGCCAGC	GCTGGGCC T GGACGATA
2114	GGCCUGU CUGAUGAG X CGAA IAUUUCGU	ACGATATC C ACAGGGCC
2115	AGGCCUG CUGAUGAG X CGAA IGAUUCG	CGATATCC A CAGGGCCT
2117	CCAGGCC CUGAUGAG X CGAA IUGGAUUA	ATATCCAC A GGGCCTGG
2122	GUGCGCCA CUGAUGAG X CGAA ICCUGUG	CACAGGGC C TGGCGCAC
2123	GGUGCGCC CUGAUGAG X CGAA IGCCUGU	ACAGGGCC T GCGCACCC
2129	CACGAAGG CUGAUGAG X CGAA ICGCCAGG	CCTGGCGC A CCTTCGTG
2131	AGCACGAA CUGAUGAG X CGAA IUGCGCCA	TGGCGCAC C TTCGTGCT
2132	CAGCACGA CUGAUGAG X CGAA IGUGCGCC	GGCGCACCC T TCGTGCTG
2139	GCACACGC CUGAUGAG X CGAA ICACGAAG	CTTCGTGC T GCGTGTGC
2152	GGGUCCUG CUGAUGAG X CGAA ICCCGCAC	GTGCGGGC C CAGGACCC
2153	CGGUCCU CUGAUGAG X CGAA IGCCCGCA	TGCGGGCC C AGGACCCG

Table 14

2154	GCGGGUCC CUGAUGAG X CGAA IGGCCCGC	GCGGGCCC A GGACCCGC
2159	AGGCGGCG CUGAUGAG X CGAA IUCCUGGG	CCCAGGAC C CGCCGCTT
2160	CAGGCGGC CUGAUGAG X CGAA IGUCCUGG	CCAGGACC C GCCGCTTG
2163	GCUCAGGC CUGAUGAG X CGAA ICGGGUCC	GGACCCGC C GCCTGAGC
2166	ACAGCUCA CUGAUGAG X CGAA ICGGCGGG	CCCGCCGC C TGAGCTGT
2167	UACAGCUC CUGAUGAG X CGAA ICGGCGCG	CCGCGGCC T GAGCTGTA
2172	CAAAGUAC CUGAUGAG X CGAA ICUCAGGC	GCCTGAGC T GTACTTTG
2177	CUUGACAA CUGAUGAG X CGAA IUACAGCU	AGCTGTAC T TTGTCAAG
2183	AUCCACCU CUGAUGAG X CGAA IACAAAGU	ACTTTGTC A AGGTGGAT
2210	GGGGAUGG CUGAUGAG X CGAA IUCGUACG	CGTACGAC A CCATCCCC
2212	UGGGGAU CUGAUGAG X CGAA IUGUCGUA	TACGACAC C ATCCCCCA
2213	CUGGGGGA CUGAUGAG X CGAA IGUGUCGU	ACGACACC A TCCCCCAG
2216	GUCCUGGG CUGAUGAG X CGAA IAUGGUGU	ACACCATC C CCCAGGAC
2217	UGUCCUGG CUGAUGAG X CGAA IGAUGGUG	CACCATCC C CCAGGACA
2218	CUGUCCUG CUGAUGAG X CGAA IGGAUGGU	ACCATCCC C CAGGACAG
2219	CCUGUCCU CUGAUGAG X CGAA IGGGAUGG	CCATCCCC C AGGACAGG
2220	GCCUGUCC CUGAUGAG X CGAA IGGGAUG	CATCCCC A GGACAGGC
2225	CGUGAGCC CUGAUGAG X CGAA IUCCUGGG	CCCAGGAC A GGCTCACG
2229	CCUCCGUG CUGAUGAG X CGAA ICCUGUCC	GGACAGGC T CACGGAGG
2231	GACCUCCG CUGAUGAG X CGAA IAGCCUGU	ACAGGCTC A CGGAGGTC
2240	GCUGGCGA CUGAUGAG X CGAA IACCUCCG	CGGAGGTC A TCGCCAGC
2245	AUGAUGCU CUGAUGAG X CGAA ICGAUGAC	GTCATCGC C AGCATCAT
2246	GAUGAUGC CUGAUGAG X CGAA IGCGAUGA	TCATCGCC A GCATCATC
2249	UUUGAUGA CUGAUGAG X CGAA ICUGGCGA	TCGCCAGC A TCATCAAA
2252	GGGUUGA CUGAUGAG X CGAA IAUGCUGG	CCAGCATC A TCAAACCC
2255	CUGGGGUU CUGAUGAG X CGAA IAUGAUGC	GCATCATC A AACCCAG
2259	UGUUCUGG CUGAUGAG X CGAA IUUGAUG	CATCAAAC C CCAGAACA
2260	GUGUUCUG CUGAUGAG X CGAA IGUUGAU	ATCAAACC C CAGAACAC
2261	CGUGUUCU CUGAUGAG X CGAA IGGUUGA	TCAAACCC C AGAACACG
2262	ACGUGUUC CUGAUGAG X CGAA IGGGUUUG	CAAACCCC A GAACACGT
2267	GCAGUACG CUGAUGAG X CGAA IUUCUGGG	CCCAGAAC A CGTACTGC
2273	ACGCACGC CUGAUGAG X CGAA IUACGUGU	ACACGTAC T CGTGCCT
2290	UGGACCAC CUGAUGAG X CGAA ICAUACCG	CGGTATGC C GTGGTCCA
2297	GGCCUUCU CUGAUGAG X CGAA IACCACGG	CCGTGGTC C AGAAGGCC
2298	CGGCCUUC CUGAUGAG X CGAA IGACCACG	CGTGGTCC A GAAGGCCG
2305	CCAUGGGC CUGAUGAG X CGAA ICCUUCUG	CAGAAGGC C GCCCATGG
2308	UGCCCAUG CUGAUGAG X CGAA ICGCCUUC	AAGGCCGC C CATGGGCA
2309	GUGCCCAU CUGAUGAG X CGAA ICGGCCU	AGGCCGCC C ATGGGCAC
2310	CGUGCCCA CUGAUGAG X CGAA IGGCGGCC	GGCCGCC A TGGGCACG
2316	UGCGGACG CUGAUGAG X CGAA ICCAUGG	CCATGGGC A CGTCCGCA
2321	GGCCUUGC CUGAUGAG X CGAA IACGUGCC	GGCACGTC C GCAAGGCC
2324	GAAGGCCU CUGAUGAG X CGAA ICGGACGU	ACGTCCGC A AGGCCTTC
2329	CUCUUGAA CUGAUGAG X CGAA ICCUUGCG	CGCAAGGC C TTCAAGAG
2330	GCUCUUGA CUGAUGAG X CGAA IGCCUUGC	GCAAGGCC T TCAAGAGC
2333	GUGGCUCU CUGAUGAG X CGAA IAAGGCCU	AGGCCTTC A AGAGCCAC
2339	AGAGACGU CUGAUGAG X CGAA ICUCUUGA	TCAAGAGC C ACGTCTCT
2340	UAGAGACG CUGAUGAG X CGAA IGCUCUUG	CAAGAGCC A CGTCTCTA

Table 14

2345	CAAGGUAG CUGAUGAG X CGAA IACGUGGC	GCCACGTC T CTACCTTG
2347	GUCAAGGU CUGAUGAG X CGAA IAGACGUG	CACGTCTC T ACCTTGAC
2350	UCUGUCAA CUGAUGAG X CGAA IUAGAGAC	GTCTCTAC C TTGACAGA
2351	GUCUGUCA CUGAUGAG X CGAA IGUAGAGA	TCTCTACC T TGACAGAC
2356	UGGAGGUC CUGAUGAG X CGAA IUCAAGGU	ACCTTGAC A GACCTCCA
2360	CGGCUGGA CUGAUGAG X CGAA IUCUGUCA	TGACAGAC C TCCAGCCG
2361	ACGGCUGG CUGAUGAG X CGAA IGUCUGUC	GACAGACC T CCAGCCGT
2363	GUACGGCU CUGAUGAG X CGAA IAGGUCUG	CAGACCTC C AGCCGTAC
2364	UGUACGGC CUGAUGAG X CGAA IGAGGUCU	AGACCTCC A GCCGTACA
2367	GCAUGUAC CUGAUGAG X CGAA ICUGGAGG	CCTCCAGC C GTACATGC
2372	CUGUCGCA CUGAUGAG X CGAA IUACGGCU	AGCCGTAC A TCGACAG
2379	CCACGAAC CUGAUGAG X CGAA IUCGCAUG	CATGCGAC A GTTCGTGG
2389	UGCAGGUG CUGAUGAG X CGAA ICCACGAA	TTCGTGGC T CACCTGCA
2391	CCUGCAGG CUGAUGAG X CGAA IAGCCACG	CGTGGCTC A CCTGCAGG
2393	CUCCUGCA CUGAUGAG X CGAA IUGAGCCA	TGGCTCAC C TGCAAGGAG
2394	UCUCCUGC CUGAUGAG X CGAA IGUGAGCC	GGCTCACC T GCAGGAGA
2397	UGGUCUCC CUGAUGAG X CGAA ICAGGUGA	TCACCTGC A GGAGACCA
2404	AGCGGGCU CUGAUGAG X CGAA IUCUCCUG	CAGGAGAC C AGCCCGCT
2405	CAGCGGGC CUGAUGAG X CGAA IGUCUCCU	AGGAGACC A GCCCGCTG
2408	CCUCAGCG CUGAUGAG X CGAA ICUGGUCU	AGACCAGC C CGCTGAGG
2409	CCCUCAGC CUGAUGAG X CGAA IGCUGGUC	GACCAGCC C GCTGAGGG
2412	CAUCCCTC CUGAUGAG X CGAA ICGGGCUG	CAGCCCGC T GAGGGATG
2422	AUGACGAC CUGAUGAG X CGAA ICAUCCCU	AGGGATGC C GTCGTCAT
2429	CUGCUCGA CUGAUGAG X CGAA IACGACGG	CCGTCGTC A TCGAGCAG
2436	AGGAGCUC CUGAUGAG X CGAA ICUCGAUG	CATCGAGC A GAGCTCCT
2441	CAGGGAGG CUGAUGAG X CGAA ICUCUGCU	AGCAGAGC T CCTCCCTG
2443	UUCAGGGA CUGAUGAG X CGAA IAGCUCUG	CAGAGCTC C TCCCTGAA
2444	AUUCAGGG CUGAUGAG X CGAA IGAGCUCU	AGAGCTCC T CCCTGAAT
2446	UCAUUCAG CUGAUGAG X CGAA IAGGAGCU	AGCTCCTC C CTGAATGA
2447	CUCAUUCA CUGAUGAG X CGAA IAGGAGC	GCTCCTCC C TGAATGAG
2448	CCUCAUUC CUGAUGAG X CGAA IGGAGGAG	CTCCTCCC T GAATGAGG
2458	CCACUGCU CUGAUGAG X CGAA ICCUCAU	AATGAGGC C AGCAGTGG
2459	GCCACUGC CUGAUGAG X CGAA IGCCUCAU	ATGAGGCC A GCAGTGGC
2462	GAGGCCAC CUGAUGAG X CGAA ICUGGCCU	AGGCCAGC A GTGGCCTC
2468	GUCCAAGA CUGAUGAG X CGAA ICCACUGC	GCAGTGGC C TCTCGAC
2469	CGUCGAAG CUGAUGAG X CGAA IGCCACUG	CAGTGGCC T CTTCGACG
2471	GACGUCGA CUGAUGAG X CGAA IAGGCCAC	GTGGCCTC T TCGACGTC
2480	GCGUAGGA CUGAUGAG X CGAA IACGUCGA	TCGACGTC T TCCTACGC
2483	GAAGCGUA CUGAUGAG X CGAA IAAGACGU	ACGTCTTC C TACGCTTC
2484	UGAAGCGU CUGAUGAG X CGAA IGAAGACG	CGTCTTCC T ACGCTTCA
2489	GCACAUGA CUGAUGAG X CGAA ICGUAGGA	TCCTACGC T TCATGTGC
2492	GUGGCACA CUGAUGAG X CGAA IAAGCGUA	TACGCTTC A TGTGCCAC
2498	GGCGUGGU CUGAUGAG X CGAA ICACAUGA	TCATGTGC C ACCACGCC
2499	CGGCGUGG CUGAUGAG X CGAA IGCACAUG	CATGTGCC A CCACGCCG
2501	CACGGCGU CUGAUGAG X CGAA IUGGCACA	TGTGCCAC C ACGCCGTG
2502	GCACGGCG CUGAUGAG X CGAA IGUGGCAC	GTGCCACC A CGCCGTGC
2506	AUGCGCAC CUGAUGAG X CGAA ICGUGGUG	CACCACGC C GTGCGCAT

Table 14

2513	GCCCCUGA CUGAUGAG X CGAA ICGCACGG		CCGTGCGC A TCAGGGGC	
2516	CUUGCCCC CUGAUGAG X CGAA IAUGCGCA		TGCGCATC A GGGGCAAG	
2522	GUAGGACU CUGAUGAG X CGAA ICCCUUGA		TCAGGGGC A AGTCCTAC	
2527	UGGACGUA CUGAUGAG X CGAA IACUUGCC		GGCAAGTC C TACGTCCA	
2528	CUGGACGU CUGAUGAG X CGAA IGACUUGC		GCAAGTCC T ACGTCCAG	
2534	CUGGCACU CUGAUGAG X CGAA IACGUAGG		CCTACGTC C AGTGCCAG	
2535	CCUGGCAC CUGAUGAG X CGAA IGACGUAG		CTACGTCC A GTGCCAGG	
2540	GAUCCCCU CUGAUGAG X CGAA ICACUGGA		TCCAGTGC C AGGGGATC	
2541	GGAUCCCC CUGAUGAG X CGAA IGCACUGG		CCAGTGCC A GGGGATCC	
2549	GCCCUGCG CUGAUGAG X CGAA IAUCCCCU		AGGGGATC C CGCAGGGC	
2550	AGCCCUGC CUGAUGAG X CGAA IGAUCCCC		GGGGATCC C GCAGGGCT	
2553	UGGAGCCC CUGAUGAG X CGAA ICGGGAUC		GATCCCGC A GGGCTCCA	
2558	GAGGAUGG CUGAUGAG X CGAA ICCUGCG		CGCAGGGC T CCATCCTC	
2560	GAGAGGAU CUGAUGAG X CGAA IAGCCUG		CAGGGCTC C ATCTCTC	
2561	GGAGAGGA CUGAUGAG X CGAA IGAGCCCU		AGGGCTCC A TCCTCTCC	
2564	CGUGGAGA CUGAUGAG X CGAA IAUGGAGC		GCTCCATC C TCTCCACG	
2565	GCGUGGAG CUGAUGAG X CGAA IGAUGGAG		CTCCATCC T CTCCACGC	
2567	CAGCGUGG CUGAUGAG X CGAA IAGGAUGG		CCATCCTC T CCACGCTG	
2569	AGCAGCGU CUGAUGAG X CGAA IAGAGGAU		ATCTCTC C ACGCTGCT	
2570	GAGCAGCG CUGAUGAG X CGAA IGAGAGGA		TCCTCTCC A CGCTGCTC	
2574	UGCAGAGC CUGAUGAG X CGAA ICGUGGAG		CTCCACGC T GCTCTGCA	
2577	GGCUGCAG CUGAUGAG X CGAA ICAGCGUG		CACGCTGC T CTGCAGCC	
2579	CAGGCUGC CUGAUGAG X CGAA IAGCAGCG		CGCTGCTC T GCAGCCTG	
2582	GCACAGGC CUGAUGAG X CGAA ICAGAGCA		TGCTCTGC A GCCTGTGC	
2585	GUAGCACA CUGAUGAG X CGAA ICUGCAGA		TCTGCAGC C TGTGCTAC	
2586	CGUAGCAC CUGAUGAG X CGAA IGCUGCAG		CTGCAGCC T GTGCTACG	
2591	GUCGCCGU CUGAUGAG X CGAA ICACAGGC		GCCTGTGC T ACGGCGAC	
2600	GUUCUCCA CUGAUGAG X CGAA IUCCCGU		ACGGCGAC A TGGAGAAC	
2609	AAACAGCU CUGAUGAG X CGAA IUUCUCCA		TGGAGAAC A AGCTGTTT	
2613	CCGCAAAC CUGAUGAG X CGAA ICUGUUC		GAACAAGC T GTTTGCGG	
2640	GCAGGAGC CUGAUGAG X CGAA ICCCGUCC		GGACGGGC T GCTCCTGC	
2643	AACGCAGG CUGAUGAG X CGAA ICAGCCCG		CGGGCTGC T CCTGCGTT	
2645	CAAACGCA CUGAUGAG X CGAA IAGCAGCC		GGCTGCTC C TCGTTTG	
2646	CCAAACGC CUGAUGAG X CGAA IGAGCAGC		GCTGCTCC T GCGTTTGG	
2666	CACCAACA CUGAUGAG X CGAA IAAAUCAU		ATGATTTC T TGTGTTG	
2677	AGGUGAGG CUGAUGAG X CGAA IUCACCAA		TTGGTGAC A CCTCACCT	
2679	UGAGGUGA CUGAUGAG X CGAA IUGUCACC		GGTGACAC C TCACCTCA	
2680	GUGAGGUG CUGAUGAG X CGAA IGUGUCAC		GTGACACC T CACCTCAC	
2682	GGGUGAGG CUGAUGAG X CGAA IAGGUGUC		GACACCTC A CCTCACCC	
2684	GUGGGUGA CUGAUGAG X CGAA IUGAGGUG		CACCTCAC C TCACCCAC	
2685	CGUGGGUG CUGAUGAG X CGAA IGUGAGGU		ACCTCACC T CACCCACG	
2687	CGCGUGGG CUGAUGAG X CGAA IAGGUGAG		CTCACCTC A CCCACCG	
2689	UUCGCGUG CUGAUGAG X CGAA IUGAGGUG		CACCTCAC C CACGCGAA	
2690	UUUCGCGU CUGAUGAG X CGAA IGUGAGGU		ACCTCACC C ACGCGAAA	
2691	UUUUCGCG CUGAUGAG X CGAA IGGUGAGG		CCTCACCC A CGCGAAA	
2701	CUGAGGAA CUGAUGAG X CGAA IUUUUCGC		GCGAAAAC C TTCCTCAG	
2702	CCUGAGGA CUGAUGAG X CGAA IGUUUUCG		CGAAAACC T TCCTCAGG	

Table 14

2705	GGUCCUGA CUGAUGAG X CGAA IAAGGUUU		AAACCTTC C TCAGGACC	
2706	GGGUCCUG CUGAUGAG X CGAA IGAAGGUU		AACCTTCC T CAGGACCC	
2708	CAGGGUCC CUGAUGAG X CGAA IAGGAAGG		CCTTCCTC A GGACCCTG	
2713	CGGACCAG CUGAUGAG X CGAA IUCCUGAG		CTCAGGAC C CTGGTCCG	
2714	UCGGACCA CUGAUGAG X CGAA IGUCCUGA		TCAGGACC C TGGTCCGA	
2715	CUCGGACC CUGAUGAG X CGAA IGGUCCUG		CAGGACCC T GGTCCGAG	
2720	GACACCUC CUGAUGAG X CGAA IACCAGGG		CCCTGGTC C GAGGTGTC	
2729	AUACUCAG CUGAUGAG X CGAA IACACCUC		GAGGTGTC C CTGAGTAT	
2730	CAUACUCA CUGAUGAG X CGAA IGACACCU		AGGTGTCC C TGAGTATG	
2731	CCAUACUC CUGAUGAG X CGAA IGGACACC		GGTGTCCC T GAGTATGG	
2741	CACCACGC CUGAUGAG X CGAA ICCAUACU		AGTATGGC T GCGTGGTG	
2753	CUUCCGCA CUGAUGAG X CGAA IUUCACCA		TGGTGAAC T TGCGGAAG	
2764	UUCACCAC CUGAUGAG X CGAA IUCUCCG		CGGAAGAC A GTGGTGAA	
2774	UACAGGGA CUGAUGAG X CGAA IUUCACCA		TGGTGAAC T TCCCTGTA	
2777	UUCUACAG CUGAUGAG X CGAA IAAGUUCA		TGAACCTC C CTGTAGAA	
2778	CUUCUACA CUGAUGAG X CGAA IGAAGUUC		GAACCTCC C TGTAAGA	
2779	UCUUCUAC CUGAUGAG X CGAA IGGAAGUU		AACTTCCC T GTAGAAGA	
2794	CCACCCAG CUGAUGAG X CGAA ICCUCGUC		GACGAGGC C CTGGGTGG	
2795	GCCACCCA CUGAUGAG X CGAA IGCCUCGU		ACGAGGCC C TGGGTGGC	
2796	UGCCACCC CUGAUGAG X CGAA IGGCCUCG		CGAGGCC T GGGTGGCA	
2804	AAAAGCCG CUGAUGAG X CGAA ICCACCCA		TGGGTGGC A CGGCTTTT	
2809	UGAACAAA CUGAUGAG X CGAA ICCUGGCC		GGCACGGC T TTTGTTCA	
2817	CCGGCAUC CUGAUGAG X CGAA IAACAAAA		TTTTGTTC A GATGCCGG	
2823	CGUGGGCC CUGAUGAG X CGAA ICAUCUGA		TCAGATGC C GGCCACG	
2827	AGGCCGUG CUGAUGAG X CGAA ICCGGCAU		ATGCCGGC C CACGGCCT	
2828	UAGGCCGU CUGAUGAG X CGAA IGCCGGCA		TGCCGGCC C ACGGCCTA	
2829	AUAGGCCG CUGAUGAG X CGAA IGGCCGGC		GCCGGCCC A CGGCCTAT	
2834	GGGGAUA CUGAUGAG X CGAA ICCUGGG		CCCACGGC C TATTCCCC	
2835	AGGGGAU CUGAUGAG X CGAA IGCCUGGG		CCACGGCC T ATTCCCCT	
2840	GCACCAGG CUGAUGAG X CGAA IAAUAGGC		GCCTATTC C CTTGGTGC	
2841	CGCACCAG CUGAUGAG X CGAA IGAUAGG		CCTATTCC C CTGGTGCG	
2842	CCGCACCA CUGAUGAG X CGAA IGGAUAG		CTATTCCC C TGGTGCGG	
2843	GCCGCACC CUGAUGAG X CGAA IGGGAUA		TATTCCCC T GGTGCGGC	
2852	CAGCAGCA CUGAUGAG X CGAA ICCGCACC		GGTGCGGC C TGCTGCTG	
2853	CCAGCAGC CUGAUGAG X CGAA IGCCGCAC		GTGCGGCC T GCTGCTGG	
2856	UAUCCAGC CUGAUGAG X CGAA ICAGGCCG		CGGCCTGC T GCTGGATA	
2859	GGGUAUCC CUGAUGAG X CGAA ICAGCAGG		CCTGCTGC T GGATACCC	
2866	AGGUCCG CUGAUGAG X CGAA IUAUCCAG		CTGGATAC C CGGACCTT	
2867	CAGGGUCC CUGAUGAG X CGAA IGUAUCCA		TGGATACC C GGACCCTG	
2872	ACCUCCAG CUGAUGAG X CGAA IUCCGGGU		ACCCGGAC C CTGGAGGT	
2873	CACCUCCA CUGAUGAG X CGAA IGUCCGGG		CCCGGACC C TGGAGGTG	
2874	GCACCUCC CUGAUGAG X CGAA IGGUCCGG		CCGGACCC T GGAGGTGC	
2883	AGUCGCUC CUGAUGAG X CGAA ICACCUCC		GGAGGTGC A GAGCGACT	
2891	GCUGGAGU CUGAUGAG X CGAA IUCGCUCU		AGAGCGAC T ACTCCAGC	
2894	AUAGCUGG CUGAUGAG X CGAA IUAGUCGC		GCGACTAC T CCAGCTAT	
2896	GCAUAGCU CUGAUGAG X CGAA IAGUAGUC		GACTACTC C AGCTATGC	
2897	GGCAUAGC CUGAUGAG X CGAA IGAGUAGU		ACTACTCC A GCTATGCC	

Table 14

2900	CCGGGCAU CUGAUGAG X CGAA ICUGGAGU	ACTCCAGC T ATGCCCGG
2905	GAGGUCCG CUGAUGAG X CGAA ICAUAGCU	AGCTATGC C CGGACCTC
2906	GGAGGUCC CUGAUGAG X CGAA IGCAUAGC	GCTATGCC C GGACCTCC
2911	CUGAUGGA CUGAUGAG X CGAA IUCCGGGC	GCCCCGAC C TCCATCAG
2912	UCUGAUGG CUGAUGAG X CGAA IGUCCGGG	CCCCGACC T CCATCAGA
2914	GCUCUGAU CUGAUGAG X CGAA IAGGUCCG	CGGACCTC C ATCAGAGC
2915	GGCUCUGA CUGAUGAG X CGAA IGAGGUCC	GGACCTCC A TCAGAGCC
2918	ACUGGCUC CUGAUGAG X CGAA IAUUGAGG	CCTCCATC A GAGCCAGT
2923	GUGAGACU CUGAUGAG X CGAA ICUCUGAU	ATCAGAGC C AGTCTCAC
2924	GGUGAGAC CUGAUGAG X CGAA IGCUCUGA	TCAGAGCC A GTCTCACC
2928	UGAAGGUG CUGAUGAG X CGAA IACUGGCU	AGCCAGTC T CACCTTCA
2930	GUUGAAGG CUGAUGAG X CGAA IAGACUGG	CCAGTCTC A CCTTCAAC
2932	CGGUUGAA CUGAUGAG X CGAA IUGAGACU	AGTCTCAC C TTCAACCG
2933	GCGGUUGA CUGAUGAG X CGAA IGUGAGAC	GTCTCACC T TCAACCGC
2936	GCCGCGGU CUGAUGAG X CGAA IAAGGUGA	TCACCTTC A ACCGCGGC
2939	GAAGCCGC CUGAUGAG X CGAA IUUGAAGG	CCTTCAAC C GCGGCTTC
2945	AGCCUUGA CUGAUGAG X CGAA ICCGCGGU	ACCGCGGC T TCAAGGCT
2948	CCCAGCCU CUGAUGAG X CGAA IAGCCGC	GCGGCTTC A AGGCTGGG
2953	UUCUCCC CUGAUGAG X CGAA ICCUUGAA	TTCAAGGC T GGGAGGAA
2963	GCGACGCA CUGAUGAG X CGAA IUUCCUCC	GGAGGAAC A TCGTTCGC
2972	AAAGAGUU CUGAUGAG X CGAA ICGACGCA	TGCGTCGC A AACTCTTT
2976	CCCCAAAG CUGAUGAG X CGAA IUUUGCGA	TCGCAAAC T CTTTGGGG
2978	GACCCCAA CUGAUGAG X CGAA IAGUUUGC	GCAAATC T TTGGGGTC
2987	CAGCCGCA CUGAUGAG X CGAA IACCCCAA	TGGGGGTC T TGCGGCTG
2994	GACACUUC CUGAUGAG X CGAA ICCGCAAG	CTTGCGGC T GAAGTGTC
3003	ACAGGCUG CUGAUGAG X CGAA IACACUUC	GAAGTGTC A CAGCCTGT
3005	AAACAGGC CUGAUGAG X CGAA IUGACACU	AGTGTAC A GCCTGTTT
3008	CAGAAACA CUGAUGAG X CGAA ICUGUGAC	GTCACAGC C TGTTCTG
3009	CCAGAAAC CUGAUGAG X CGAA IGCUGUGA	TCACAGCC T GTTCTGG
3015	GCAAAUCC CUGAUGAG X CGAA IAAACAGG	CCTGTTTC T GGATTGTC
3024	UGUUCACC CUGAUGAG X CGAA ICAAAUCC	GGATTGTC A GGTGAACA
3032	CUGGAGGC CUGAUGAG X CGAA IUUCACCU	AGGTGAAC A GCCTCCAG
3035	CGUCUGGA CUGAUGAG X CGAA ICUGUUCA	TGAACAGC C TCCAGACG
3036	CCGUCUGG CUGAUGAG X CGAA IGCUGUUC	GAACAGCC T CCAGACGG
3038	CACCGUCU CUGAUGAG X CGAA IAGGCUGU	ACAGCCTC C AGACGGTG
3039	ACACCGUC CUGAUGAG X CGAA IGAGGCUG	CAGCCTCC A GACGGTGT
3050	GAUGUUGG CUGAUGAG X CGAA ICACACCG	CGGTGTGC A CCAACATC
3052	UAGAUGUU CUGAUGAG X CGAA IUGCACAC	GTGTGCAC C AACATCTA
3053	GUAGAUGU CUGAUGAG X CGAA IGUGCACA	TGTGCACC A ACATCTAC
3056	CUUGUAGA CUGAUGAG X CGAA IUUGGUGC	GCACCAAC A TCTACAAG
3059	GAUCUUGU CUGAUGAG X CGAA IAUUGUGG	CCAACATC T ACAAGATC
3062	GAGGAUCU CUGAUGAG X CGAA IUAGAUGU	ACATCTAC A AGATCCTC
3068	CAGCAGGA CUGAUGAG X CGAA IAUUUGU	ACAAGATC C TCCTGCTG
3069	GCAGCAGG CUGAUGAG X CGAA IGAUCUUG	CAAGATCC T CCTGCTGC
3071	CUGCAGCA CUGAUGAG X CGAA IAGGAUCU	AGATCCTC C TGCTGCAG
3072	CCUGCAGC CUGAUGAG X CGAA IGAGGAUC	GATCCTCC T GCTGCAGG
3075	ACGCCUGC CUGAUGAG X CGAA ICAGGAGG	CCTCCTGC T GCAGGCGT

Table 14

3078	UGUACGCC CUGAUGAG X CGAA ICAGCAGG	CCTGCTGC A GGCCTACA
3086	GUGAAACC CUGAUGAG X CGAA IUACGCCU	AGGCGTAC A GGTTCAC
3093	CACAUGCG CUGAUGAG X CGAA IAAACCUG	CAGGTTTC A CGCATGTG
3097	AGCACACA CUGAUGAG X CGAA ICGUGAAA	TTTCACGC A TGTGTGCT
3105	GGAGCUGC CUGAUGAG X CGAA ICACACAU	ATGTGTGC T GCAGCTCC
3108	AUGGGAGC CUGAUGAG X CGAA ICAGCACA	TGTGCTGC A GCTCCCAT
3111	GAAAUGGG CUGAUGAG X CGAA ICUGCAGC	GCTGCAGC T CCCATTTC
3113	AUGAAAUG CUGAUGAG X CGAA IAGCUGCA	TGCAGCTC C CATTTCAT
3114	GAUGAAAU CUGAUGAG X CGAA IGAGCUGC	GCAGCTCC C ATTTTCATC
3115	UGAUGAAA CUGAUGAG X CGAA IGGAGCUG	CAGCTCCC A TTTTCATCA
3120	CUUGCUGA CUGAUGAG X CGAA IAAAUUGG	CCCATTTC A TCAGCAAG
3123	AAACUUGC CUGAUGAG X CGAA IAUGAAAU	ATTTTCATC A GCAAGTTT
3126	UCCAAACU CUGAUGAG X CGAA ICUGAUGA	TCATCAGC A AGTTTGA
3140	AAAUGUGG CUGAUGAG X CGAA IUUCUUC	GGAAGAAC C CCACATT
3141	AAAAUGUG CUGAUGAG X CGAA IGUUCUUC	GAAGAACC C CACATT
3142	AAAAAUGU CUGAUGAG X CGAA IGGUUCUU	AAGAACCC C ACATTTT
3143	GAAAAAUG CUGAUGAG X CGAA IGGUUCUU	AGAACCCC A CATTTTTC
3145	AGGAAAAA CUGAUGAG X CGAA IUUGGGUU	AACCCAC A TTTTCT
3152	GACGCGCA CUGAUGAG X CGAA IAAAAAUG	CATTTTTC C TGCGCGTC
3153	UGACGCGC CUGAUGAG X CGAA IGAAAAAU	ATTTTTC T GCGCGTCA
3161	GUCAGAGA CUGAUGAG X CGAA IACGCGCA	TGCGCGTC A TCTCTGAC
3164	CGUGUCAG CUGAUGAG X CGAA IAUGACGC	GCGTCATC T CTGACACG
3166	GCCGUGUC CUGAUGAG X CGAA IAGAUGAC	GTCATCTC T GACACGGC
3170	GGAGGCCG CUGAUGAG X CGAA IUCAGAGA	TCTCTGAC A CGGCCTCC
3175	CAGAGGGA CUGAUGAG X CGAA ICCGUGUC	GACACGGC C TCCCTCTG
3176	GCAGAGGG CUGAUGAG X CGAA IGCCGUGU	ACACGGCC T CCCTCTGC
3178	UAGCAGAG CUGAUGAG X CGAA IAGGCCGU	ACGGCCTC C CTCTGCTA
3179	GUAGCAGA CUGAUGAG X CGAA IGAGGCCG	CGGCCTCC C TCTGCTAC
3180	AGUAGCAG CUGAUGAG X CGAA IGGAGGCC	GGCCTCCC T CTGCTACT
3182	GGAGUAGC CUGAUGAG X CGAA IAGGGAGG	CCTCCCTC T GCTACTCC
3185	GAUGGAGU CUGAUGAG X CGAA ICAGAGGG	CCCTCTGC T ACTCCATC
3188	CAGGAUGG CUGAUGAG X CGAA IUAGCAGA	TCTGCTAC T CCATCCTG
3190	UUCAGGAU CUGAUGAG X CGAA IAGUAGCA	TGCTACTC C ATCTGAA
3191	UUUCAGGA CUGAUGAG X CGAA IGAGUAGC	GCTACTCC A TCCTGAAA
3194	GGCUUUA CUGAUGAG X CGAA IAUGGAGU	ACTCCATC C TGAAAGCC
3195	UGGCUUUC CUGAUGAG X CGAA IGAUGGAG	CTCCATCC T GAAAGCCA
3202	GCGUUCU CUGAUGAG X CGAA ICUUUCAG	CTGAAAGC C AAGAACGC
3203	UGCGUUCU CUGAUGAG X CGAA ICGUUUA	TGAAAGCC A AGAACGCA
3211	GACAUCCC CUGAUGAG X CGAA ICGUUCU	AAGAACGC A GGGATGTC
3222	UGGCCCC CUGAUGAG X CGAA ICGACAUC	GATGTCGC T GGGGGCCA
3229	GCGCCCU CUGAUGAG X CGAA ICCCCAG	CTGGGGGC C AAGGGCGC
3230	GGCGCCCU CUGAUGAG X CGAA IGCCCCA	TGGGGGCC A AGGGGCC
3238	GGGCCGGC CUGAUGAG X CGAA ICGCCCU	AAGGGCGC C GCCGGCCC
3241	AGAGGGCC CUGAUGAG X CGAA ICGGCGCC	GGCGCCGC C GGCCCTCT
3245	GGGCAGAG CUGAUGAG X CGAA ICCGGCGG	CCGCCGGC C CTCTGCCC
3246	AGGGCAGA CUGAUGAG X CGAA IGCCGGCG	CGCCGGCC C TCTGCCCT
3247	GAGGGCAG CUGAUGAG X CGAA IGGCCGGC	GCCGGCCC T CTGCCCTC

Table 14

3249	CGGAGGGC CUGAUGAG X CGAA IAGGGCCG	CGGCCCTC T GCCCTCCG
3252	CCUCGGAG CUGAUGAG X CGAA ICAGAGGG	CCCTCTGC C CTCCGAGG
3253	GCCUCGGA CUGAUGAG X CGAA IGCAGAGG	CCTCTGCC C TCCGAGGC
3254	GGCCUCGG CUGAUGAG X CGAA IGGCAGAG	CTCTGCCC T CCGAGGCC
3256	ACGGCCUC CUGAUGAG X CGAA IAGGGCAG	CTGCCCTC C GAGGCCGT
3262	CACUGCAC CUGAUGAG X CGAA ICCUCGGA	TCCGAGGC C GTGCAGTG
3267	ACAGCCAC CUGAUGAG X CGAA ICACGGCC	GGCCGTGC A GTGGCTGT
3273	GGUGGCAC CUGAUGAG X CGAA ICCACUGC	GCAGTGGC T GTGCCACC
3278	UGCUUGGU CUGAUGAG X CGAA ICACAGCC	GGCTGTGC C ACCAAGCA
3279	AUGCUUGG CUGAUGAG X CGAA IGCACAGC	GCTGTGCC A CCAAGCAT
3281	GAAUGCUU CUGAUGAG X CGAA IUGGCACA	TGTGCCAC C AAGCATTC
3282	GGAAUGCU CUGAUGAG X CGAA IGUGGCAC	GTGCCACC A AGCATTC
3286	AGCAGGAA CUGAUGAG X CGAA ICUUGGUG	CACCAAGC A TTCCTGCT
3290	CUUGAGCA CUGAUGAG X CGAA IAAUGCUU	AAGCATTC C TGCTCAAG
3291	GCUUGAGC CUGAUGAG X CGAA IGAAUGCU	AGCATTC T GCTCAAGC
3294	UCAGCUUG CUGAUGAG X CGAA ICAGGAAU	ATTCCTGC T CAAGCTGA
3296	AGUCAGCU CUGAUGAG X CGAA IAGCAGGA	TCCTGCTC A AGCTGACT
3300	GUCGAGUC CUGAUGAG X CGAA ICUUGAGC	GCTCAAGC T GACTCGAC
3304	CGGUGUCG CUGAUGAG X CGAA IUCAGCUU	AAGCTGAC T CGACACCG
3309	UGACACGG CUGAUGAG X CGAA IUCGAGUC	GACTCGAC A CCGTGTC
3311	GGUGACAC CUGAUGAG X CGAA IUGUCGAG	CTCGACAC C GTGTCACC
3317	CACGUAGG CUGAUGAG X CGAA IACACGGU	ACCGTGTC A CCTACGTG
3319	GGCACGUA CUGAUGAG X CGAA IUGACACG	CGTGTCAC C TACGTGCC
3320	UGGCACGU CUGAUGAG X CGAA IGUGACAC	GTGTCACC T ACGTGCCA
3327	CCAGGAGU CUGAUGAG X CGAA ICACGUAG	CTACGTGC C ACTCCTGG
3328	CCCAGGAG CUGAUGAG X CGAA IGCACGUA	TACGTGCC A CTCCTGGG
3330	ACCCCAGG CUGAUGAG X CGAA IUGGCACG	CGTGCCAC T CCTGGGGT
3332	UGACCCCA CUGAUGAG X CGAA IAGUGGCA	TGCCACTC C TGGGGTCA
3333	GUGACCCC CUGAUGAG X CGAA IGAGUGGC	GCCACTCC T GGGGTCAC
3340	GUCCUGAG CUGAUGAG X CGAA IACCCCAG	CTGGGGTC A CTCAGGAC
3342	CUGUCCUG CUGAUGAG X CGAA IUGACCCC	GGGGTCAC T CAGGACAG
3344	GGCUGUCC CUGAUGAG X CGAA IAGUGACC	GGTCACTC A GGACAGCC
3349	GUCUGGGC CUGAUGAG X CGAA IUCCUGAG	CTCAGGAC A GCCCAGAC
3352	UGCGUCUG CUGAUGAG X CGAA ICUGUCCU	AGGACAGC C CAGACGCA
3353	CUGCGUCU CUGAUGAG X CGAA IGCUGUCC	GGACAGCC C AGACGCAG
3354	GCUGCGUC CUGAUGAG X CGAA IGGCUGUC	GACAGCCC A GACGCAGC
3360	GACUCAGC CUGAUGAG X CGAA ICGUCUGG	CCAGACGC A GCTGAGTC
3363	UCCGACUC CUGAUGAG X CGAA ICUGCGUC	GACGCAGC T GAGTCGGA
3375	UCCCCGGG CUGAUGAG X CGAA ICUUCCGA	TCGGAAGC T CCCGGGGA
3377	CGUCCCCG CUGAUGAG X CGAA IAGCUUCC	GGAAGCTC C CGGGGACG
3378	UCGUCCCC CUGAUGAG X CGAA IGAGCUUC	GAAGCTCC C GGGGACGA
3390	GGGCAGUC CUGAUGAG X CGAA ICGUCGUC	GACGACGC T GACTGCCC
3394	UCCAGGGC CUGAUGAG X CGAA IUCAGCGU	ACGCTGAC T GCCCTGGA
3397	GCCUCCAG CUGAUGAG X CGAA ICAGUCAG	CTGACTGC C CTGGAGGC
3398	GGCCUCCA CUGAUGAG X CGAA IGCAGUCA	TGACTGCC C TGGAGGCC
3399	CGGCCUCC CUGAUGAG X CGAA IGGCAGUC	GACTGCCC T GGAGGCCG
3406	UUGGCUGC CUGAUGAG X CGAA ICCUCCAG	CTGGAGGC C GCAGCCAA

Table 14

3409	GGGUUGGC CUGAUGAG X CGAA ICGGCCUC		GAGGCCGC A GCCAACCC	
3412	GCCGGGUU CUGAUGAG X CGAA ICUGCGGC		GCCGCAGC C AACCCGGC	
3413	UGCCGGGU CUGAUGAG X CGAA IGCUGCGG		CCGCAGCC A ACCCGGCA	
3416	CAGUGCCG CUGAUGAG X CGAA IUUGGCUG		CAGCCAAC C CGGCACTG	
3417	GCAGUGCC CUGAUGAG X CGAA IGUUGGCU		AGCCAACC C GGCCTGTC	
3421	GAGGGCAG CUGAUGAG X CGAA ICCGGGUU		AACCCGGC A CTGCCCTC	
3423	CUGAGGGC CUGAUGAG X CGAA IUGCCGGG		CCCGGCAC T GCCCTCAG	
3426	AGUCUGAG CUGAUGAG X CGAA ICAGUGCC		GGCACTGC C CTCAGACT	
3427	AAGUCUGA CUGAUGAG X CGAA IGAGUGC		GCACTGCC C TCAGACTT	
3428	GAAGUCUG CUGAUGAG X CGAA IGGCAGUG		CACTGCCC T CAGACTTC	
3430	UUGAAGUC CUGAUGAG X CGAA IAGGGCAG		CTGCCCTC A GACTTCAA	
3434	GGUCUUGA CUGAUGAG X CGAA IUCUGAGG		CCTCAGAC T TCAAGACC	
3437	GAUGGUCU CUGAUGAG X CGAA IAAGUCUG		CAGACTTC A AGACCATC	
3442	UCCAGGAU CUGAUGAG X CGAA IUCUUGAA		TTCAAGAC C ATCCTGGA	
3443	GUCCAGGA CUGAUGAG X CGAA IGUCUUGA		TCAAGACC A TCCTGGAC	
3446	UCAGUCCA CUGAUGAG X CGAA IAUGGUCU		AGACCATC C TGGACTGA	
3447	AUCAGUCC CUGAUGAG X CGAA IGAUGGUC		GACCATCC T GGACTGAT	
3452	UGGCAUCC CUGAUGAG X CGAA IUCCAGGA		TCCTGGAC T GATGGCCA	
3459	GGGCGGGU CUGAUGAG X CGAA ICCAUCAG		CTGATGGC C ACCCGCCC	
3460	UGGGCGGG CUGAUGAG X CGAA IGCCAUCA		TGATGGCC A CCCGCCCA	
3462	UGUGGGCG CUGAUGAG X CGAA IUGGCCAU		ATGGCCAC C CGCCACA	
3463	CUGUGGGC CUGAUGAG X CGAA IGUGGCCA		TGGCCACC C GCCCACAG	
3466	UGGUGUGU CUGAUGAG X CGAA ICGGUGUG		CCACCCGC C CACAGCCA	
3467	CUGGUGUG CUGAUGAG X CGAA ICGGGUGU		CACCCGCC C ACAGCCAG	
3468	CCUGGCUG CUGAUGAG X CGAA IGGCGGGU		ACCGCCCC A CAGCCAGG	
3470	GGCUGGC CUGAUGAG X CGAA IUGGCGG		CCGCCAC A GCCAGGCC	
3473	CUCGGCCU CUGAUGAG X CGAA ICUGUGGG		CCCACAGC C AGGCCGAG	
3474	UCUCGGCC CUGAUGAG X CGAA IGCUGUGG		CCACAGCC A GGCCGAGA	
3478	CUGCUCUC CUGAUGAG X CGAA ICCUGGCU		AGCCAGGC C GAGAGCAG	
3485	CUGGUGUC CUGAUGAG X CGAA ICUCUCGG		CCGAGAGC A GACACCAG	
3489	GCUGCUGG CUGAUGAG X CGAA IUCUGCUC		GAGCAGAC A CCAGCAGC	
3491	GGGUGUCU CUGAUGAG X CGAA IUGUCUGC		GCAGACAC C AGCAGCCC	
3492	AGGGCUGC CUGAUGAG X CGAA IGUGUCUG		CAGACACC A GCAGCCCT	
3495	GACAGGGC CUGAUGAG X CGAA ICUGGUGU		ACACCAGC A GCCCTGTC	
3498	CGUGACAG CUGAUGAG X CGAA ICUGCUGG		CCAGCAGC C CTGTCACG	
3499	GCGUGACA CUGAUGAG X CGAA IGCUGCUG		CAGCAGCC C TGTACGCG	
3500	GGCGUGAC CUGAUGAG X CGAA IGGCUGCU		AGCAGCCC T GTCACGCC	
3504	GCCCGGCG CUGAUGAG X CGAA IACAGGGC		GCCCTGTC A CGCCGGGC	
3508	UAGAGCCC CUGAUGAG X CGAA ICGUGACA		TGTCACGC C GGGCTCTA	
3513	GGACGUAG CUGAUGAG X CGAA ICCCGGCG		CGCCGGGC T CTACGTCC	
3515	UGGGACCU CUGAUGAG X CGAA IAGCCCGG		CCGGGCTC T ACGTCCCA	
3521	CCUCCUG CUGAUGAG X CGAA IACGUAGA		TCTACGTC C CAGGAGG	
3522	CCCUCCU CUGAUGAG X CGAA IGACGUAG		CTACGTCC C AGGGAGG	
3523	UCCCUCC CUGAUGAG X CGAA IGGACGUA		TACGTCCC A GGGAGGGA	
3540	UGGGUGUG CUGAUGAG X CGAA ICCGCCCC		GGGCGGCG C CACACCCA	
3541	CUGGGUGU CUGAUGAG X CGAA IGCCGCCC		GGGCGGCG C ACACCCAG	
3542	CCUGGGUG CUGAUGAG X CGAA IGGCGGCC		GGCGGCCC A CACCCAGG	

Table 14

3544	GGCCUGGG CUGAUGAG X CGAA IUGGGCCG		CGGCCAC A CCCAGGCC	
3546	CGGGCCUG CUGAUGAG X CGAA IUGUGGGC		GCCCACAC C CAGGCCCG	
3547	GCGGGCCU CUGAUGAG X CGAA IGUGUGGG		CCCACACC C AGGCCCGC	
3548	UGCGGGCC CUGAUGAG X CGAA IGGUGUGG		CCACACCC A GGCCCGCA	
3552	GCGGUGCG CUGAUGAG X CGAA ICCUGGGU		ACCCAGGC C CGCACCGC	
3553	AGCGGUGC CUGAUGAG X CGAA IGCCUGGG		CCCAGGCC C GCACCGCT	
3556	CCCAGCGG CUGAUGAG X CGAA ICGGGCCU		AGGCCCGC A CCGCTGGG	
3558	CUCCAGC CUGAUGAG X CGAA IUGCGGGC		GCCCGCAC C GCTGGAG	
3561	AGACUCCC CUGAUGAG X CGAA ICGUGCG		CGCACCGC T GGGAGTCT	
3569	CAGGCCUC CUGAUGAG X CGAA IACUCCA		TGGGAGTC T GAGGCCTG	
3575	CUCACUCA CUGAUGAG X CGAA ICCUCAGA		TCTGAGGC C TGAGTGAG	
3576	ACUCACUC CUGAUGAG X CGAA IGCCUCAG		CTGAGGCC T GAGTGAGT	
3592	CAGGCCUC CUGAUGAG X CGAA ICCAAACA		TGTTTGGC C GAGGCCTG	
3598	GACAUGCA CUGAUGAG X CGAA ICCUCGGC		GCCGAGGC C TGCATGTC	
3599	GGACAUGC CUGAUGAG X CGAA IGCCUCGG		CCGAGGCC T GCATGTCC	
3602	GCCGGACA CUGAUGAG X CGAA ICAGGCCU		AGGCCTGC A TGTCCGGC	
3607	CUUCAGCC CUGAUGAG X CGAA IACAUGCA		TGCATGTC C GGCTGAAG	
3611	CAGCCUUC CUGAUGAG X CGAA ICCGGACA		TGTCCGGC T GAAGGCTG	
3618	GGACACUC CUGAUGAG X CGAA ICCUUCAG		CTGAAGGC T GAGTGTC	
3626	CCUCAGCC CUGAUGAG X CGAA IACACUCA		TGAGTGTC C GGCTGAGG	
3630	CAGGCCUC CUGAUGAG X CGAA ICCGGACA		TGTCCGGC T GAGGCCTG	
3636	CUCGCUCA CUGAUGAG X CGAA ICCUCAGC		GCTGAGGC C TGAGCGAG	
3637	ACUCGCUC CUGAUGAG X CGAA IGCCUCAG		CTGAGGCC T GAGCGAGT	
3649	CCUUGGCU CUGAUGAG X CGAA IACACUCG		CGAGTGTC C AGCCAAGG	
3650	CCCUUGGC CUGAUGAG X CGAA IGACACUC		GAGTGTC A GCCAAGGG	
3653	CAGCCCUU CUGAUGAG X CGAA ICUGGACA		TGTCCAGC C AAGGGCTG	
3654	UCAGCCCU CUGAUGAG X CGAA IGCUGGAC		GTCCAGCC A AGGGCTGA	
3660	GGACACUC CUGAUGAG X CGAA ICCUUGG		CCAAGGGC T GAGTGTC	
3668	GGUGUGCU CUGAUGAG X CGAA IACACUCA		TGAGTGTC C AGCACACC	
3669	AGGUGUGC CUGAUGAG X CGAA IGACACUC		GAGTGTC A GCACACCT	
3672	GGCAGGUG CUGAUGAG X CGAA ICUGGACA		TGTCCAGC A CACCTGCC	
3674	ACGGCAGG CUGAUGAG X CGAA IUGUGGA		TCCAGCAC A CCTGCCGT	
3676	AGACGGCA CUGAUGAG X CGAA IUGUGCUG		CAGCACAC C TGCGTCT	
3677	AAGACGGC CUGAUGAG X CGAA IGUGUGCU		AGCACACC T GCGTCTT	
3680	GUGAAGAC CUGAUGAG X CGAA ICAGGUGU		ACACCTGC C GTCTTCAC	
3684	GGAAGUGA CUGAUGAG X CGAA IACGGCAG		CTGCCGTC T TCACTTCC	
3687	UGGGGAAG CUGAUGAG X CGAA IAAGACGG		CCGTCTTC A CTCCCCA	
3689	UGUGGGGA CUGAUGAG X CGAA IUGAAGAC		GTCTTCAC T TCCCCA	
3692	GCCUGUGG CUGAUGAG X CGAA IAAGUGAA		TCACTTC C CCACAGGC	
3693	AGCCUGUG CUGAUGAG X CGAA IGAAGUGA		TCACTTCC C CACAGGCT	
3694	CAGCCUGU CUGAUGAG X CGAA IGGAAGUG		CACTTCCC C ACAGGCTG	
3695	CCAGCCUG CUGAUGAG X CGAA IGGGAAGU		ACTTCCCC A CAGGCTGG	
3697	CGCCAGCC CUGAUGAG X CGAA IUGGGGAA		TTCCCCAC A GGCTGGCG	
3701	CGAGCGCC CUGAUGAG X CGAA ICCUGUGG		CCACAGGC T GGCGCTCG	
3707	UGGAGCCG CUGAUGAG X CGAA ICGCCAGC		GCTGGCGC T CGGCTCCA	
3712	UGGGGUGG CUGAUGAG X CGAA ICCGAGCG		CGCTCGGC T CCACCCCA	
3714	CCUGGGGU CUGAUGAG X CGAA IAGCCGAG		CTCGGCTC C ACCCCAGG	

Table 14

3715	CCCUGGGG CUGAUGAG X CGAA IGAGCCGA	TCGGCTCC A CCCAGGG
3717	GGCCCUGG CUGAUGAG X CGAA IUGGAGCC	GGCTCCAC C CCAGGGCC
3718	UGGCCUG CUGAUGAG X CGAA IGUGGAGC	GCTCCACC C CAGGGCCA
3719	CUGGCCCU CUGAUGAG X CGAA IGGUGGAG	CTCCACCC C AGGCCAG
3720	GCUGGCC CUGAUGAG X CGAA IGGUGGA	TCCACCCC A GGGCCAGC
3725	GAAAAGCU CUGAUGAG X CGAA ICCUGGG	CCCAGGGC C AGCTTTTC
3726	GGAAAAGC CUGAUGAG X CGAA IGCCCUGG	CCAGGGCC A GCTTTTCC
3729	UGAGGAAA CUGAUGAG X CGAA ICUGGCC	GGGCCAGC T TTTCTCA
3734	CCUGGUGA CUGAUGAG X CGAA IAAAAGCU	AGCTTTTC C TCACCAGG
3735	UCCUGGUG CUGAUGAG X CGAA IGAAAAGC	GCTTTTCC T CACCAGGA
3737	GCUCCUGG CUGAUGAG X CGAA IAGGAAAA	TTTTCTC A CCAGGAGC
3739	GGGCUCCU CUGAUGAG X CGAA IUGAGGAA	TTCTCAC C AGGAGCCC
3740	CGGGCUCC CUGAUGAG X CGAA IGUGAGGA	TCCTACC A GGAGCCCG
3746	GGAAGCCG CUGAUGAG X CGAA ICUCCUGG	CCAGGAGC C CGGCTTCC
3747	UGGAAGCC CUGAUGAG X CGAA IGCUCUG	CAGGAGCC C GGCTTCCA
3751	GGAGUGGA CUGAUGAG X CGAA ICCGGGCU	AGCCCGGC T TCCACTCC
3754	UGGGGAGU CUGAUGAG X CGAA IAAGCCGG	CCGGCTC C ACTCCCCA
3755	GUGGGGAG CUGAUGAG X CGAA IGAAGCCG	CGGCTCC A CTCCCCAC
3757	AUGUGGGG CUGAUGAG X CGAA IUUGAAGC	GCTTCCAC T CCCACAT
3759	CUAUGUGG CUGAUGAG X CGAA IAGUGGAA	TTCCACTC C CCACATAG
3760	CCUAUGUG CUGAUGAG X CGAA IGAGUGGA	TCCACTCC C CACATAGG
3761	UCCUAUGU CUGAUGAG X CGAA IGGAGUGG	CCACTCCC C ACATAGGA
3762	UUCCUAUG CUGAUGAG X CGAA IGGGAGUG	CACTCCCC A CATAGGAA
3764	UAUUCUA CUGAUGAG X CGAA IUGGGGAG	CTCCCCAC A TAGGAATA
3776	CUGGGGAU CUGAUGAG X CGAA IACUAUUC	GAATAGTC C ATCCCCAG
3777	UCUGGGGA CUGAUGAG X CGAA IGACUAUU	AATAGTCC A TCCCCAGA
3780	GAAUCUGG CUGAUGAG X CGAA IAUGGACU	AGTCCATC C CCAGATTC
3781	CGAAUCUG CUGAUGAG X CGAA IGAUGGAC	GTCCATCC C CAGATTCC
3782	GCGAAUCU CUGAUGAG X CGAA IGGAUGGA	TCCATCCC C AGATTCCG
3783	GGCGAAUC CUGAUGAG X CGAA IGGGAUGG	CCATCCCC A GATTCCGC
3791	UGAACAAU CUGAUGAG X CGAA ICGAAUCU	AGATTCCG C ATTGTTC
3792	GUGAACAA CUGAUGAG X CGAA IGCGAAUC	GATTCCGC A TTGTTCAC
3799	GCGAGGGG CUGAUGAG X CGAA IAACAAUG	CATTGTTC A CCCCTCGC
3801	GGGCGAGG CUGAUGAG X CGAA IUGAACAA	TTGTTCAC C CCTCGCCC
3802	AGGGCGAG CUGAUGAG X CGAA IGUGAAC	TGTTCACC C CTCGCCCT
3803	CAGGGCGA CUGAUGAG X CGAA IGGUGAAC	GTTACCCC C TCGCCCTG
3804	GCAGGGCG CUGAUGAG X CGAA IGGUGAA	TTACCCCC T CGCCCTGC
3808	GAGGGCAG CUGAUGAG X CGAA ICGAGGGG	CCCCTCGC C CTGCCCTC
3809	GGAGGGCA CUGAUGAG X CGAA ICGAGGGG	CCCTCGCC C TGCCCTCC
3810	AGGAGGGC CUGAUGAG X CGAA IGGCGAGG	CCTCGCCC T GCCCTCCT
3813	CAAAGGAG CUGAUGAG X CGAA ICAGGGCG	CGCCCTGC C CTCTTTG
3814	GCAAAGGA CUGAUGAG X CGAA ICGAGGGC	GCCCTGCC C TCCTTTGC
3815	GGCAAAGG CUGAUGAG X CGAA IGGCAGGG	CCCTGCCC T CCTTTGCC
3817	AAGGCAAA CUGAUGAG X CGAA IAGGGCAG	CTGCCCTC C TTTGCCTT
3818	GAAGGCAA CUGAUGAG X CGAA IGAGGGCA	TGCCCTCC T TTGCCTT
3823	GGGUGGAA CUGAUGAG X CGAA ICAAAGGA	TCCTTTGC C TTCCACCC
3824	GGGUGGA CUGAUGAG X CGAA IGCAAAGG	CCTTTGCC T TCCACCCC

Table 14

3827	GUGGGGGU CUGAUGAG X CGAA IAAGGCCAA	TTGCCTTC C ACCCCCAC
3828	GGUGGGGG CUGAUGAG X CGAA IGAAGGCA	TGCCTTCC A CCCCCACC
3830	AUGGUGGG CUGAUGAG X CGAA IUGGAAGG	CCTTCCAC C CCCACCAT
3831	GAUGGUGG CUGAUGAG X CGAA IGUGGAAG	CTTCCACC C CCACCATC
3832	GGAUGGUG CUGAUGAG X CGAA IGGUGGAA	TTCCACCC C CACCATCC
3833	UGGAUGGU CUGAUGAG X CGAA IGGGUGGA	TCCACCCC C ACCATCCA
3834	CUGGAUGG CUGAUGAG X CGAA IGGGUGGG	CCACCCCC A CCATCCAG
3836	ACCUGGAU CUGAUGAG X CGAA IUGGGGGU	ACCCCCAC C ATCCAGGT
3837	CACCUGGA CUGAUGAG X CGAA IUGGGGGG	CCCCCACC A TCCAGGTG
3840	CUCCACCU CUGAUGAG X CGAA IAUGGUGG	CCACCATC C AGGTGGAG
3841	UCUCCACC CUGAUGAG X CGAA IGAUGGUG	CACCATCC A GGTGGAGA
3851	CUUCUCAG CUGAUGAG X CGAA IUCUCCAC	GTGGAGAC C CTGAGAAG
3852	CCUUCUCA CUGAUGAG X CGAA IGUCUCCA	TGGAGACC C TGAGAAGG
3853	UCCUUCUC CUGAUGAG X CGAA IGGUCUCC	GGAGACCC T GAGAAGGA
3863	GCUCCCAG CUGAUGAG X CGAA IUCCUUCU	AGAAGGAC C CTGGGAGC
3864	AGCUCCCA CUGAUGAG X CGAA IGUCCUUC	GAAGGACC C TGGGAGCT
3865	GAGCUCCC CUGAUGAG X CGAA IGGUCCUU	AAGGACCC T GGGAGCTC
3872	AUUCCCAG CUGAUGAG X CGAA ICUCCCAG	CTGGGAGC T CTGGGAAT
3874	AAAUUCCC CUGAUGAG X CGAA IAGCUCCC	GGGAGCTC T GGGAAATT
3891	ACACCUUU CUGAUGAG X CGAA IUCACUCC	GGAGTGAC C AAAGGTGT
3892	CACACCUU CUGAUGAG X CGAA IGUCACUC	GAGTGACC A AAGGTGTG
3902	GUGUACAG CUGAUGAG X CGAA ICACACCU	AGGTGTGC C CTGTACAC
3903	UGUGUACA CUGAUGAG X CGAA IGCACACC	GGTGTGCC C GTACACA
3904	CUGUGUAC CUGAUGAG X CGAA IGGCACAC	GTGTGCCC T GTACACAG
3909	CUCGCCUG CUGAUGAG X CGAA IUACAGGG	CCCTGTAC A CAGGCGAG
3911	UCCUCGCC CUGAUGAG X CGAA IUGUACAG	CTGTACAC A GCGAGGA
3921	AGGUGCAG CUGAUGAG X CGAA IUCCUCGC	GCGAGGAC C CTGCACCT
3922	CAGGUGCA CUGAUGAG X CGAA IGUCCUCG	CGAGGACC C TGCACCTG
3923	CCAGGUGC CUGAUGAG X CGAA IGGUCCUC	GAGGACCC T GCACCTGG
3926	CAUCCAGG CUGAUGAG X CGAA ICAGGGUC	GACCCTGC A CCTGGATG
3928	CCCAUCCA CUGAUGAG X CGAA IUGCAGGG	CCCTGCAC C TGGATGGG
3929	CCCCAUCC CUGAUGAG X CGAA IGUGCAGG	CCTGCACC T GGATGGGG
3941	ACCCACAG CUGAUGAG X CGAA IACCCCCA	TGGGGGTC C CTGTGGGT
3942	GACCCACA CUGAUGAG X CGAA IGACCCCC	GGGGGTCC C TGTGGGTC
3943	UGACCCAC CUGAUGAG X CGAA IGGACCCC	GGGGTCCC T GTGGGTCA
3951	CCCCAAUU CUGAUGAG X CGAA IACCCACA	TGTGGGTC A AATTGGGG
3968	ACUCCAC CUGAUGAG X CGAA ICACCUCC	GGAGGTGC T GTGGGAGT
3984	AUAUAUUC CUGAUGAG X CGAA IUAUUUUA	TAAAATAC T GAATATAT
4002	UUAUAAAC CUGAUGAG X CGAA IAAAAACU	AGTTTTTC A GTTTTGAA

Stem Length = 8 . Core Sequence = CUGAUGAG X CGAA (X = GCCGUUAGGC or other stem II sequence and length (greater than or equal to 2 base-pairs)). I = Inosine nucleotide

Seq1 = TERT (Homo sapiens telomerase reverse transcriptase (TERT) mRNA, 4015 bp); Nakamura *et al.*, Science 277 (5328), 955-959 (1997)

Table 15

Table 15: Human telomerase reverse transcriptase (TERT) G-Cleaver Ribozyme and Target Sequence

nt. Position	Substrate Sequence	Seq ID Nos	Ribozyme Sequence	Seq ID Nos
16	GCUGGUGCU G CUGCG		CGCAG UGAUGGCAUGCACAUGCGCG AGGACGCAGC	
19	GGGUCCUGCU G CGCAC		GUGCG UGAUGGCAUGCACAUGCGCG AGCAGGACGC	
21	GUCCUGUGC G CACGU		ACGUG UGAUGGCAUGCACAUGCGCG GCAGCAGGAC	
53	GGCCACCCCG G CGAUG		CAUCG UGAUGGCAUGCACAUGCGCG GGGGGUGGCC	
55	CCACCCCGCG G AUGCC		GGCAU UGAUGGCAUGCACAUGCGCG GCGGGGUGG	
58	CCCCCGGGAU G CCGCG		CGCGG UGAUGGCAUGCACAUGCGCG AUGCGGGGG	
61	CCGCGAUGCC G CGCGC		CGCGG UGAUGGCAUGCACAUGCGCG GGCAUCGCGG	
63	GGGAUGCCCG G CGCUC		GAGCG UGAUGGCAUGCACAUGCGCG GCGGCAUCGC	
65	GAUGCCGCG G CUCCC		GGGAG UGAUGGCAUGCACAUGCGCG GCGCGGCAUC	
72	CGGCUCCCG G CUGCC		GGCAG UGAUGGCAUGCACAUGCGCG GGGGAGCGCG	
75	GUUCCCGCG G CCGAG		CUCGG UGAUGGCAUGCACAUGCGCG AGCGGGGAGC	
78	CCCCGUGCC G AGCCG		CGGCU UGAUGGCAUGCACAUGCGCG GGCAGCGGGG	
85	GGCGAGCCGU G CGCUC		GAGCG UGAUGGCAUGCACAUGCGCG ACGGCUCCGC	
87	CGAGCCGUG G CUCCC		GGGAG UGAUGGCAUGCACAUGCGCG GCACGGCUCG	
94	UGCGCUCCCU G CUGCG		CGCAG UGAUGGCAUGCACAUGCGCG AGGGAGCGCA	
97	GUCCCGUGCU G CGCAG		CUGCG UGAUGGCAUGCACAUGCGCG AGCAGGGAGC	
99	UCCUGUGCG G CAGCC		GGCUG UGAUGGCAUGCACAUGCGCG GCAGCAGGGA	
111	AGCCACUACC G CGAGG		CCUCG UGAUGGCAUGCACAUGCGCG GGUAGUGGU	
113	CCACUACCG G AGGUG		CACCU UGAUGGCAUGCACAUGCGCG GCGGUAGUGG	
118	ACCGCGAGGU G CUGCC		GGCAG UGAUGGCAUGCACAUGCGCG ACCUCGCGGU	
121	GGGAGGUGCU G CCGCU		AGCGG UGAUGGCAUGCACAUGCGCG AGCACCUCGC	
124	AGGUGCUGCC G CUGGC		GCCAG UGAUGGCAUGCACAUGCGCG GGCAGCACCU	
139	CCACGUUCGU G CGGCG		CGCCG UGAUGGCAUGCACAUGCGCG ACGAACGUGG	
144	UUCGUGCGGC G CCUGG		CCAGG UGAUGGCAUGCACAUGCGCG GCCGCACGAA	
172	GGCGGUGGU G CAGCG		CGCUG UGAUGGCAUGCACAUGCGCG ACCAGCCGCC	
177	CUGGUGCAGC G CGGGG		CCCCG UGAUGGCAUGCACAUGCGCG GCUCCACCCAG	
198	GGGCUUUCG G CGCGC		GCGCG UGAUGGCAUGCACAUGCGCG GGAAGCCCGC	
200	GGCUUUCGCG G CGCUG		CAGCG UGAUGGCAUGCACAUGCGCG GCGGAAAGCC	

Table 15

202	CUUUCGCGC G CUGGU		ACCAG UGAUGGCAUGCACUAUGCGCG GCGCGGAAG	
216	GUGGCCAGU G CCUGG		CCAGG UGAUGGCAUGCACUAUGCGCG ACUGGGCCAC	
223	AGUGCCUGGU G UGCGU		ACGCA UGAUGGCAUGCACUAUGCGCG ACCAGGCACU	
225	UGCCUGGUGU G CGUGC		GCACG UGAUGGCAUGCACUAUGCGCG ACACCAGGCA	
229	UGGUGUGGU G CCCUG		CAGGG UGAUGGCAUGCACUAUGCGCG ACGCACACCA	
239	GCCUGGGAC G CACGG		CCGUG UGAUGGCAUGCACUAUGCGCG GUCCACAGGC	
247	ACGCACGGCC G CCCCC		GGGGG UGAUGGCAUGCACUAUGCGCG GGCCTGUGGU	
254	GCGCCCCC G CCGCC		GGCGG UGAUGGCAUGCACUAUGCGCG GGGGGGCGC	
257	GCCCCCGCC G CCCCC		GGGGG UGAUGGCAUGCACUAUGCGCG GGCGGGGGC	
270	CCUCCUUC G CCAGG		CCUGG UGAUGGCAUGCACUAUGCGCG GGAAGGAGGG	
277	UCCGCCAGGU G UCCUG		CAGGA UGAUGGCAUGCACUAUGCGCG ACCUGGGGA	
282	CAGGUGUCCU G CCUGA		UCAGG UGAUGGCAUGCACUAUGCGCG AGGACACCUG	
286	UGUCCUGCCU G AGGA		UCCUU UGAUGGCAUGCACUAUGCGCG AGGCAGGACA	
303	CUGGUGGCC G AGUGC		GCACU UGAUGGCAUGCACUAUGCGCG GGGCCACCAG	
307	UGGCCCGAGU G CUGCA		UGCAG UGAUGGCAUGCACUAUGCGCG ACUGGGGCA	
310	CCGAGUGCU G CAGAG		CUCUG UGAUGGCAUGCACUAUGCGCG AGCACUCGGG	
319	UGCAGAGGU G UGCGA		UCGCA UGAUGGCAUGCACUAUGCGCG AGCCUCUGCA	
321	CAGAGGCU G CGAGC		GCUCG UGAUGGCAUGCACUAUGCGCG ACAGCCUCUG	
323	GAGGCUUGC G AGCGC		GCGCU UGAUGGCAUGCACUAUGCGCG GCACAGCCUC	
327	CUGUGGAGC G CCGCG		CGCCG UGAUGGCAUGCACUAUGCGCG GCUCGCUCG	
332	CGAGCGGCG G CGAAG		CUUCG UGAUGGCAUGCACUAUGCGCG GCGCGCGCU	
334	AGCGCGGCG G AAGAA		UUCUU UGAUGGCAUGCACUAUGCGCG GCGCGCGCU	
343	CGAAGAACGU G CUGGC		GCCAG UGAUGGCAUGCACUAUGCGCG ACGUUCUUCG	
359	CUUCGCUUC G CGCUG		CAGCG UGAUGGCAUGCACUAUGCGCG GAAGCCGAAG	
361	UCGGCUUCGC G CUGCU		AGCAG UGAUGGCAUGCACUAUGCGCG GCGAAGCCGA	
364	GCUUCGCGU G CUGGA		UCCAG UGAUGGCAUGCACUAUGCGCG AGCGGAAGC	
378	GACGGGCCC G CCGGG		CCCCG UGAUGGCAUGCACUAUGCGCG GGGCCCCGUC	
392	GGCCCCCCC G AGGCC		GGCCU UGAUGGCAUGCACUAUGCGCG GGGGGGGCCC	
412	CCACGAGGU G CGCAG		CUGCG UGAUGGCAUGCACUAUGCGCG ACGCUGGUGG	
414	ACCAGGUGC G CAGCU		AGCUG UGAUGGCAUGCACUAUGCGCG GCACGCUUGU	
424	GCAGCUACCU G CCGAA		UUGGG UGAUGGCAUGCACUAUGCGCG AGGUAGCUCG	

Table 15

436	CAAACACGGU G ACCGA	UCCGU UGAUGGCAUGCACUAUGCGCG ACCGUGUUGG	
440	CACGGUACC G AGCA	UCCGU UGAUGGCAUGCACUAUGCGCG GGUCACCGUG	
443	GGUACCGAC G CACUG	CAGUG UGAUGGCAUGCACUAUGCGCG GUCGGUCACC	
448	CCGACGCACU G CGGGG	CCCCG UGAUGGCAUGCACUAUGCGCG AGUGCGUCGG	
472	CGUGGGGCU G CUGCU	AGCAG UGAUGGCAUGCACUAUGCGCG AGCCCCACG	
475	GGGGGCU G CUGCG	CGCAG UGAUGGCAUGCACUAUGCGCG AGCAGCCCCC	
478	GGCUGCUGCU G CGCGG	CGGCG UGAUGGCAUGCACUAUGCGCG AGCAGCAGCC	
480	CUGCUGCUGC G CGCGG	CGCGG UGAUGGCAUGCACUAUGCGCG GCAGCAGCAG	
483	CUGCUGCGCC G CGUGG	CCACG UGAUGGCAUGCACUAUGCGCG GGCGCAGCAG	
491	CCGCGUGGGC G ACGAC	GUCGU UGAUGGCAUGCACUAUGCGCG GCCCAGCGCG	
494	CGUGGGCGAC G ACGUG	CACGU UGAUGGCAUGCACUAUGCGCG GUCGCCACCG	
499	GCACGACGU G CUGGU	ACCAG UGAUGGCAUGCACUAUGCGCG ACGUCGUCGC	
511	UGGUUACCU G CUGGC	GCCAG UGAUGGCAUGCACUAUGCGCG AGGUGAACCA	
519	CUGCUGGCAC G CUGCG	CGCAG UGAUGGCAUGCACUAUGCGCG GUGCCAGCAG	
522	CUGGCACGU G CGCGC	GCGCG UGAUGGCAUGCACUAUGCGCG AGCGUGCCAG	
524	GGCACGCUGC G CGCUC	GAGCG UGAUGGCAUGCACUAUGCGCG GCAGCGUGCC	
526	CACGCUGGC G CUCUU	AAGAG UGAUGGCAUGCACUAUGCGCG GCGCAGCGUG	
533	CGCGCUCUU G UGCUG	CAGCA UGAUGGCAUGCACUAUGCGCG AAAGAGCGCG	
535	CGCUCUUUGU G CUGGU	ACCAG UGAUGGCAUGCACUAUGCGCG ACAGAGAGCG	
552	GUCCCCAGCU G CGCCU	AGGCG UGAUGGCAUGCACUAUGCGCG AGCUGGGAGC	
554	UCCACGUCG G CCUAC	GUAGG UGAUGGCAUGCACUAUGCGCG GCAGCUGGGA	
565	CCUACCAAGU G UGC GG	CCGCA UGAUGGCAUGCACUAUGCGCG ACCUGGUAGG	
567	UACCAAGGUGU G CGGGC	GCCCC UGAUGGCAUGCACUAUGCGCG ACACUUGGUA	
574	UGUGCGGGCC G CGGCU	AGCGG UGAUGGCAUGCACUAUGCGCG GGCCCCGACA	
577	GCGGGCCGCC G CUGUA	UACAG UGAUGGCAUGCACUAUGCGCG GCGGGCCCGC	
580	GGCGGCCGCU G UACCA	UGGUA UGAUGGCAUGCACUAUGCGCG AGCGGGCGCC	
593	CCAGCUCGSC G CUGCC	GGCAG UGAUGGCAUGCACUAUGCGCG GCGGAGCUGG	
596	GUUGGGGCGU G CCACU	AGUGG UGAUGGCAUGCACUAUGCGCG AGCGCCGAGC	
616	CCCGGCCCCC G CCACA	UGUGG UGAUGGCAUGCACUAUGCGCG GGGGGCCGGG	
623	CCGCGCACAC G CUAGU	ACUAG UGAUGGCAUGCACUAUGCGCG GUGUGGCGGG	
636	AGUGGACCCC G AAGGC	GCCUU UGAUGGCAUGCACUAUGCGCG GGGGUCCACU	

Table 13

651	CGUCUGGAU G CGAAC		GUUCG UGAUGGCAUGCACUAUGCGG AUCCAGACG	
653	UCUGGAUGC G AACGG		CCGUU UGAUGGCAUGCACUAUGCGG GCAUCCAGA	
703	CCUGGGCCU G CCAGC		GCUGG UGAUGGCAUGCACUAUGCGG AGGCCAGGG	
716	AGCCCGGGU G CGAGG		CCUCG UGAUGGCAUGCACUAUGCGG ACCCGGGCU	
718	CCCCGGGUC G AGGAG		CUCCU UGAUGGCAUGCACUAUGCGG GCACCCGGG	
726	GCGAGGAGC G CGGGG		CCCCG UGAUGGCAUGCACUAUGCGG GCCUCCUCG	
737	CGGGGCAGU G CCAGC		GCUGG UGAUGGCAUGCACUAUGCGG ACUGCCCCG	
744	AGUGCCAGC G AAGUC		GACUU UGAUGGCAUGCACUAUGCGG GGCUGGCACU	
751	GCCGAAGUCU G CCGUU		AACGG UGAUGGCAUGCACUAUGCGG AGACUUCGG	
757	GUCUGCCGUU G CCCAA		UUGGG UGAUGGCAUGCACUAUGCGG AACGGCAGC	
779	CAGGCGUGG G CUGCC		GGCAG UGAUGGCAUGCACUAUGCGG GCCACGCCUG	
782	GCGUGGCGU G CCCC		AGGGG UGAUGGCAUGCACUAUGCGG AGCGCCACG	
788	CGCUGCCCU G AGCCG		CGGCU UGAUGGCAUGCACUAUGCGG AGGGGACGG	
802	CGGAGCGAC G CCCGU		ACGGG UGAUGGCAUGCACUAUGCGG GUCCGCUCCG	
841	CGGGCAGGAC G CGUGG		CCACG UGAUGGCAUGCACUAUGCGG GUCCUGCCG	
850	CGCGUGGACC G AGUGA		UCACU UGAUGGCAUGCACUAUGCGG GGUCCACCG	
854	UGGACCGAGU G ACCGU		ACGGU UGAUGGCAUGCACUAUGCGG ACUCGGUCCA	
867	CGUGGUUUCU G UGUGG		CCACA UGAUGGCAUGCACUAUGCGG AGAAACCACG	
869	UGGUUUCUGU G UGGUG		CACCA UGAUGGCAUGCACUAUGCGG ACAGAAACCA	
874	UCUGUGUGGU G UCACC		GGUGA UGAUGGCAUGCACUAUGCGG ACCACACAGA	
881	GGUGUCACCU G CCAGA		UCUGG UGAUGGCAUGCACUAUGCGG AGGUGACACC	
890	UGCCAGACCC G CCGAA		UUCGG UGAUGGCAUGCACUAUGCGG GGGUCUGGCA	
893	CAGACCCGCC G AAGAA		UUCUU UGAUGGCAUGCACUAUGCGG GGCGGGUCUG	
917	UTUGGAGGGU G CGCUC		GAGCG UGAUGGCAUGCACUAUGCGG ACCCUCCAA	
919	UGGAGGGUGC G CUCUC		GAGAG UGAUGGCAUGCACUAUGCGG GCACCCUCCA	
931	UCUCUGGCAC G CGCCA		UGGCG UGAUGGCAUGCACUAUGCGG GUGCCAGAGA	
933	UCUGGCACGC G CCACU		AGUGG UGAUGGCAUGCACUAUGCGG GCGUGCCAGA	
957	UCCUGGGGC G CCAGC		GCUGG UGAUGGCAUGCACUAUGCGG GGCACACGGA	
968	CCAGCACCCAC G CGGGC		GCCCC UGAUGGCAUGCACUAUGCGG GUGGUGCUGG	
988	CAUCCACAUC G CGGCC		GGCCG UGAUGGCAUGCACUAUGCGG GAUGUGGAUG	
1012	CCUGGGACAC G CCUUG		CAAGG UGAUGGCAUGCACUAUGCGG GUGUCCACG	

Table 15

1017	GACACGCCUU G UCCCC		GGGGA UGAUGGCAUGCACUAUGCGCG AGGCGUGUC
1027	GUCCCCGGGU G UACGC		GGUA UGAUGGCAUGCACUAUGCGCG ACCGGGGAC
1031	CCCGUGUAC G CCGAG		CUCGG UGAUGGCAUGCACUAUGCGCG GUACACCGGG
1034	GGUGUACGCC G AGACC		GGUCU UGAUGGCAUGCACUAUGCGCG GCGGUACACC
1064	CUCCUCAGGC G ACNAG		CUUGU UGAUGGCAUGCACUAUGCGCG GCCUGAGGAG
1078	AGGAGCAGCU G CGGCC		GGCGG UGAUGGCAUGCACUAUGCGCG AGCUGCUCCU
1105	UCAGCUCUCU G AGGCC		GGCCU UGAUGGCAUGCACUAUGCGCG AGAGAGTGA
1117	GGCCACGCCU G ACUGG		CCAGU UGAUGGCAUGCACUAUGCGCG AGGUGGGCC
1124	CCUGACUGGC G CUCGG		CCGAG UGAUGGCAUGCACUAUGCGCG GCCAGUCAGG
1171	GGCCUUGGAU G CCAGG		CCUGG UGAUGGCAUGCACUAUGCGCG AUCCAGGGCC
1185	GGGACUCCCC G CAGGU		ACCUG UGAUGGCAUGCACUAUGCGCG GGGGAGUCCC
1192	CCCGCAGGUU G CCCCC		CGGGG UGAUGGCAUGCACUAUGCGCG AACUUGCGGG
1197	AGGUUGCCCC G CCUGC		GCAGG UGAUGGCAUGCACUAUGCGCG GGGGCAACCU
1201	UGCCCCGCCU G CCCCCA		UGGGG UGAUGGCAUGCACUAUGCGCG AGGCGGGCA
1209	CUGCCCCAGC G CUACU		AGUAG UGAUGGCAUGCACUAUGCGCG GCUUGGGCAG
1222	ACUGGCAAAU G CGGCC		GGCCG UGAUGGCAUGCACUAUGCGCG AUUUGCCAGU
1231	UGCGGCCCCU G UUUCU		AGAAA UGAUGGCAUGCACUAUGCGCG AGGGGCCGCA
1243	UUUGGAGCU G CUUGG		CCAAG UGAUGGCAUGCACUAUGCGCG AGCUCAGAA
1256	UGGGAACAC G CGCAG		CUGCG UGAUGGCAUGCACUAUGCGCG GUUGUCCCA
1258	GGAACACGC G CAGUG		CACUG UGAUGGCAUGCACUAUGCGCG GCGUGGUUCC
1263	CACGCGCAGU G CCCCCU		AGGGG UGAUGGCAUGCACUAUGCGCG ACUGCGCGUG
1276	CCUACGGGUU G CUCCU		AGGAG UGAUGGCAUGCACUAUGCGCG ACCCCGUAGG
1288	UCCUCAAGAC G CACUG		CAGUG UGAUGGCAUGCACUAUGCGCG GUCUUGAGGA
1293	AAGACGCACU G CCGGC		GCGGG UGAUGGCAUGCACUAUGCGCG AGUGCGUCUU
1297	CGCACUGCCC G CUGCG		CGCAG UGAUGGCAUGCACUAUGCGCG GGGCAGUGCG
1300	ACUGCCCGCU G CGAGC		GCUCG UGAUGGCAUGCACUAUGCGCG AGCGGGCAGU
1302	UGCCCGCUGC G AGCUG		CAGCU UGAUGGCAUGCACUAUGCGCG GCAGCGGGCA
1307	GUUGCGAGCU G CCGUC		GACCG UGAUGGCAUGCACUAUGCGCG AGCUCGCAGC
1328	AGCAGCGGUU G UCUUG		ACAGA UGAUGGCAUGCACUAUGCGCG ACCGGCUGCU
1332	GCCGGUGUCU G UGCCC		GGGCA UGAUGGCAUGCACUAUGCGCG AGACACCGGC
1334	CGGUGUCUGU G CCGCG		CCGGG UGAUGGCAUGCACUAUGCGCG ACAGACACCG

Table 15

1358	CCAGGCUUCU G UGGCG	CGCCA UGAUGGCAUGCACUAUGCGG AGAGCCUUGG
1370	GGCGCCCC G AGGAG	CUCCU UGAUGGCAUGCACUAUGCGG GGGGCCGCC
1395	GACCCCGUC G CCUGG	CCAGG UGAUGGCAUGCACUAUGCGG GACGGGGUC
1402	GUCCGCUUGU G CAGCU	AGCUG UGAUGGCAUGCACUAUGCGG ACCAGGCGAC
1408	UGGUGCAGCU G CUCCG	CGGAG UGAUGGCAUGCACUAUGCGG AGCUGCACCA
1413	CAGCUGUCC G CCAGC	GCUGG UGAUGGCAUGCACUAUGCGG GGAGCAGCU
1438	CCUGGCAGGU G UACGG	CCGUA UGAUGGCAUGCACUAUGCGG ACCUGCCAGG
1450	ACGGCUTUGU G CGGGC	GCCCG UGAUGGCAUGCACUAUGCGG ACGAAGCCGU
1458	GUGCGGGCCU G CCUGC	GCAGG UGAUGGCAUGCACUAUGCGG AGGCCCGCAC
1462	GGGCCUGCCU G CGCCG	CGGCG UGAUGGCAUGCACUAUGCGG AGGCAGGCC
1464	GCCUGCCUGC G CCGGC	GCCCG UGAUGGCAUGCACUAUGCGG GCAGGCAGGC
1474	GCCGGCUGGU G CCCCC	GGGGG UGAUGGCAUGCACUAUGCGG ACCAGCCGGC
1505	CAGGCACAAC G AACGC	GCGUU UGAUGGCAUGCACUAUGCGG GUTUGGCCUG
1509	CACAACGAAC G CCGCU	AGCGG UGAUGGCAUGCACUAUGCGG GUTCGTUGUG
1512	AACGAACGCC G CUUCC	GGAAG UGAUGGCAUGCACUAUGCGG GCGTUGGUU
1556	GGGGAAGCAU G CCAAG	CUUUG UGAUGGCAUGCACUAUGCGG AUGCUUCCCC
1567	CCAAGCUCUC G CUGCA	UGCAG UGAUGGCAUGCACUAUGCGG GAGAGCUUGG
1570	AGCUCUCGU G CAGGA	UCCUG UGAUGGCAUGCACUAUGCGG AGCGAGAGCU
1579	UGCAGGAGCU G ACGUG	CACGU UGAUGGCAUGCACUAUGCGG AGCUCUUGCA
1591	CGUGGAAGAU G AGCGU	ACGCU UGAUGGCAUGCACUAUGCGG AUCUUCACG
1597	AGAUGAGCGU G CGGGA	UCCCG UGAUGGCAUGCACUAUGCGG ACGCUCAUCU
1605	GUGCGGGACU G CCGUU	AAGCG UGAUGGCAUGCACUAUGCGG AGUCCCGCAC
1607	GCGGGACUGC G CUUGG	CCAAG UGAUGGCAUGCACUAUGCGG GCAGUCCCGC
1615	GCGCUTUGGU G CGCAG	CUGCG UGAUGGCAUGCACUAUGCGG AGCCAAGGCG
1617	GCUUGGCUGC G CAGGA	UCCUG UGAUGGCAUGCACUAUGCGG GCAGCCAAAGC
1638	GGGUTUGGU G UGUUC	GAACA UGAUGGCAUGCACUAUGCGG AGCCAACCCC
1640	GGUUGGCGUU G UUGCG	CGGAA UGAUGGCAUGCACUAUGCGG ACAGCCAAACC
1649	UGUCCCGGCC G CAGAG	CUCUG UGAUGGCAUGCACUAUGCGG GGCCGGAACA
1663	AGCACCGUCU G CGUGA	UCACG UGAUGGCAUGCACUAUGCGG AGACGGUGCU
1667	CCGUCUGCGU G AGGAG	CUCCU UGAUGGCAUGCACUAUGCGG ACGCAGACGG
1690	CCAAGUTUCCU G CACUG	CAGUG UGAUGGCAUGCACUAUGCGG AGGAACUUGG

Table 15

1699	UGCACUGGCU G AUGAG		CUCAU UGAUGGCAUGCACUAUGCGCG AGCCAGUGCA	
1702	ACUGGCUGAU G AGUGU		ACACU UGAUGGCAUGCACUAUGCGCG AUGAGCCAGU	
1706	GCUGAUGAGU G UGUAC		GUACA UGAUGGCAUGCACUAUGCGCG ACUCAUCAGC	
1708	UGAUGAGUGU G UACGU		ACGUA UGAUGGCAUGCACUAUGCGCG ACACUCAUCA	
1718	GUACGUGGUC G AGCUG		CAGCU UGAUGGCAUGCACUAUGCGCG GACGACGUAC	
1723	UCGUCGAGCU G CUCAG		CUGAG UGAUGGCAUGCACUAUGCGCG AGCUCGACGA	
1742	UUUCUUAU G UCACG		CGUGA UGAUGGCAUGCACUAUGCGCG AUAAAAGAAA	
1793	CCGGAAGAGU G UCUGG		CCAGA UGAUGGCAUGCACUAUGCGCG ACUCUCCGG	
1807	GGAGCAAGUU G CAAAG		CUTUG UGAUGGCAUGCACUAUGCGCG AACUUGCUCC	
1834	GACAGCACUU G AAGAG		CUCUU UGAUGGCAUGCACUAUGCGCG AAGUGCUGUC	
1843	UGAAGAGGGU G CAGCU		AGCUG UGAUGGCAUGCACUAUGCGCG ACCCUCUACA	
1849	GGGUGCAGCU G CGGGA		UCCCG UGAUGGCAUGCACUAUGCGCG AGCUGCACCC	
1858	UGCGGGAGCU G UCGGA		UCCGA UGAUGGCAUGCACUAUGCGCG AGCUCGCCGA	
1898	AGCCAGGCC G CCCUG		CAGGG UGAUGGCAUGCACUAUGCGCG GGGCCUUGGCU	
1903	GGCCCGCCU G CUGAC		GUCAG UGAUGGCAUGCACUAUGCGCG AGGGCGGGCC	
1906	CGGCCUGCU G ACGUC		GACGU UGAUGGCAUGCACUAUGCGCG AGCAGGGCGG	
1920	UCCAGACUCC G CUUCA		UGAAG UGAUGGCAUGCACUAUGCGCG GGAGUCUGGA	
1937	CCCCAAGCCU G ACGGG		CCCCU UGAUGGCAUGCACUAUGCGCG AGGCUUGGGG	
1945	CUGACGGGCU G CGGCC		GGCCG UGAUGGCAUGCACUAUGCGCG AGCCCCGUCAG	
1951	GGCUGCGGC G AUTUG		ACAAU UGAUGGCAUGCACUAUGCGCG GGCCGCGAGCC	
1955	GCGCCGGAU G UGAAC		GUUCA UGAUGGCAUGCACUAUGCGCG AAUCGGCCGC	
1957	GGCCGAUUG G AACAU		AUGUU UGAUGGCAUGCACUAUGCGCG ACAAUCCGCC	
1992	AGAACGUUCC G CAGAG		CUCUG UGAUGGCAUGCACUAUGCGCG GGAACGUUUCU	
2009	AAAGAGGCC G AGCGU		ACGCU UGAUGGCAUGCACUAUGCGCG GGCCCCUCUUU	
2023	GUUCACCCUC G AGGGU		ACCCU UGAUGGCAUGCACUAUGCGCG GAGGUGAGAC	
2029	CCUGAGGGU G AAGGC		GCCUU UGAUGGCAUGCACUAUGCGCG ACCCUCGAGG	
2038	UGAAGGCACU G UUCAG		CUGAA UGAUGGCAUGCACUAUGCGCG AGUGCCUUA	
2047	UGUUCAGCGU G CUCAA		UUGAG UGAUGGCAUGCACUAUGCGCG ACGCUGAACA	
2057	GCUCAACUAC G AGCGG		CCGCU UGAUGGCAUGCACUAUGCGCG GUAGUUGAGC	
2065	ACGAGCGGGC G CGGCG		CGCCG UGAUGGCAUGCACUAUGCGCG GCCCGCUCGU	
2070	CGGGCGCGGC G CCCC		CGGGG UGAUGGCAUGCACUAUGCGCG GCCCGGCCCG	

Table 15

2087	CCUCCUGGGC G CCUCU	AGAGG UGAUGGCAUGCACUAUGCGCG GCCCAGGAGG
2093	GGCGCCUCU G UGCUG	CAGCA UGAUGGCAUGCACUAUGCGCG AGAGGCGCCC
2095	GGCGCCUCU G CUGGG	CCCAG UGAUGGCAUGCACUAUGCGCG ACAGAGGCGC
2108	GGCGCCUGGAC G AUAUC	GAUUA UGAUGGCAUGCACUAUGCGCG GUCCAGGCGCC
2127	AGGGCCUGGC G CACCU	AGGUG UGAUGGCAUGCACUAUGCGCG GCCAGGCCCU
2137	GCACCUUCGU G CUGCG	CGCAG UGAUGGCAUGCACUAUGCGCG ACGAAGGUGC
2140	CCUUGGUGU G CGUGU	ACACG UGAUGGCAUGCACUAUGCGCG AGCAGGAAGG
2144	CGUGCUGGU G UGCGG	CCGCA UGAUGGCAUGCACUAUGCGCG ACGCAGCAGC
2146	UGCUGCGUGU G CGGGC	GCCCG UGAUGGCAUGCACUAUGCGCG ACACGCAGCA
2161	CCCAGGACC G CCGCC	GGCGG UGAUGGCAUGCACUAUGCGCG GGGUCCUGGG
2164	AGGACCCGCC G CUUGA	UCAGG UGAUGGCAUGCACUAUGCGCG GGGCGGUCCU
2168	CCCGCCGCCU G AGCUG	CAGCU UGAUGGCAUGCACUAUGCGCG AGGCGGCGGG
2173	CGCCUGAGCU G UACUU	AAGUA UGAUGGCAUGCACUAUGCGCG AGCUCAGGCG
2180	GCUGUACUUU G UCAAG	CUUGA UGAUGGCAUGCACUAUGCGCG AAGUACACAG
2192	CAAGGUGAU G UGACG	CGUCA UGAUGGCAUGCACUAUGCGCG AUCCACCUCG
2194	AGGUGGAUGU G ACGGG	CCCCU UGAUGGCAUGCACUAUGCGCG ACAUCCACCU
2201	UGUGACGGGC G CGUAC	GUACG UGAUGGCAUGCACUAUGCGCG GCCGUCACA
2207	GGCGCGUAC G ACACC	GGUGU UGAUGGCAUGCACUAUGCGCG GUACGCGCCC
2243	GGAGGUCAUC G CCAGC	GCUGG UGAUGGCAUGCACUAUGCGCG GAUGACCCUCC
2274	AACACGUACU G CGUGC	GCACG UGAUGGCAUGCACUAUGCGCG AGUACGGUGU
2278	CGUACUGCGU G CGUCG	CGACG UGAUGGCAUGCACUAUGCGCG ACGCAGUACG
2288	GGCUGGGUAC G CCGUG	CACGG UGAUGGCAUGCACUAUGCGCG AUACCGACGC
2306	CCAGAAGGCC G CCCAU	AUGGG UGAUGGCAUGCACUAUGCGCG GGCCTUCUGG
2322	GGGCACGUCC G CAAGG	CCUUG UGAUGGCAUGCACUAUGCGCG GGACGUGCCCC
2353	UCUCUACCUU G ACAGA	UCUGU UGAUGGCAUGCACUAUGCGCG AAGGUAGAGA
2374	AGCCGUACAU G CGACA	UGUCG UGAUGGCAUGCACUAUGCGCG AUGUACGGCU
2376	CCGUACAUGC G ACAGU	ACUGU UGAUGGCAUGCACUAUGCGCG GCAUGUACGG
2395	UGGCUACCU G CAGGA	UCCUG UGAUGGCAUGCACUAUGCGCG AGGUGAGCCA
2410	AGACCAAGCCC G CUGAG	CUCAG UGAUGGCAUGCACUAUGCGCG GGGCUGGUCU
2413	CCAGCCCGCU G AGGGA	UCCCU UGAUGGCAUGCACUAUGCGCG AGCGGCGUGG
2420	GCUGAGGGAU G CCGUC	GACGG UGAUGGCAUGCACUAUGCGCG AUCCUCACAGC

Table 15

2432	CGUCGUCAUC G AGCAG		CUGCU UGAUGGCAUGCACUAUGCGCG GAUGACGACG
2449	GCUCUCCCU G AAUGA		UCAUU UGAUGGCAUGCACUAUGCGCG AGGGAGGAGC
2453	CUCCUGAAU G AGGCC		GGCCU UGAUGGCAUGCACUAUGCGCG AUUCAGGGAG
2474	UGGCCUCUUC G ACGUC		GACGU UGAUGGCAUGCACUAUGCGCG GAAGAGGCCA
2487	GUCUUCUAC G CUUCA		UGAAG UGAUGGCAUGCACUAUGCGCG GUAGGAAGAC
2494	UACGCUUCAU G UGCCA		UGGCA UGAUGGCAUGCACUAUGCGCG AUGAAGSGUA
2496	CGCUCUAUGU G CCACC		GGUGG UGAUGGCAUGCACUAUGCGCG ACAUGAAGCG
2504	GUGCCACCAC G CCGUG		CACGG UGAUGGCAUGCACUAUGCGCG GUGGUGGCAC
2509	ACCACGCCGU G CGCAU		AUGCG UGAUGGCAUGCACUAUGCGCG ACGGCGUGGU
2511	CAGGCCGUGC G CAUCA		UGAUG UGAUGGCAUGCACUAUGCGCG GCACGGCGUG
2538	UACGUCCAGU G CCAGG		CCUGG UGAUGGCAUGCACUAUGCGCG ACUGGACGUA
2551	AGGGGAUCCC G CAGGG		CCCUG UGAUGGCAUGCACUAUGCGCG GGGAUCCCCU
2572	UCCUCUCCAC G CUGCU		AGCAG UGAUGGCAUGCACUAUGCGCG GUGGAGAGGA
2575	UCUCCACGCU G CUCUG		CAGAG UGAUGGCAUGCACUAUGCGCG AGCGUGGAGA
2580	ACGCUGCUCU G CAGCC		GGCUG UGAUGGCAUGCACUAUGCGCG AGAGCAGCGU
2587	UCUGCAGCCU G UGCUA		UAGCA UGAUGGCAUGCACUAUGCGCG AGGCUGCAGA
2589	UGCAGCCUGU G CUACG		CGUAG UGAUGGCAUGCACUAUGCGCG ACAGGCUGCA
2597	GUGCUACGGC G ACAUG		CAUGU UGAUGGCAUGCACUAUGCGCG GCCGUAGCAC
2614	AGAACAAAGCU G UUUGC		GCAAA UGAUGGCAUGCACUAUGCGCG AGCUUGUUCU
2618	CAAGCUGUUU G CGGGG		CCCCG UGAUGGCAUGCACUAUGCGCG AAACAGCUIUG
2641	GGGACGGGCU G CUCCU		AGGAG UGAUGGCAUGCACUAUGCGCG AGCCCGUCCC
2647	GGCUGCUCCU G CGUUU		AAACG UGAUGGCAUGCACUAUGCGCG AGGAGCAGCC
2660	UUUGGUGGAU G AUUUC		GAUUU UGAUGGCAUGCACUAUGCGCG AUCCACCAAA
2668	AUGAUUUUUU G UUUGU		ACCAA UGAUGGCAUGCACUAUGCGCG AAGAAUUCAU
2674	UCUUGUUGGU G ACACC		GGUGU UGAUGGCAUGCACUAUGCGCG ACCAACAGA
2693	CCUCACCCAC G CGAAA		UUUUCG UGAUGGCAUGCACUAUGCGCG GUGGGUGAGG
2695	UCACCCACGC G AAAC		GUUUU UGAUGGCAUGCACUAUGCGCG GCGUGGGUGA
2721	ACCCUGGUCC G AGGUG		CACCU UGAUGGCAUGCACUAUGCGCG GGACCAGGGU
2726	GGUCCGAGGU G UCCCU		AGGGA UGAUGGCAUGCACUAUGCGCG ACCUCGGACC
2732	AGGUGUCCCU G AGUAU		AUACU UGAUGGCAUGCACUAUGCGCG AGGGACACCU
2742	GAGUAUGGCU G CGUGG		CCACG UGAUGGCAUGCACUAUGCGCG AGCCAUACUC

Table 15

2749	GCUGCGUGGU G AACUU		AAGUU UGAUGGCAUGCACUAUGCGCG ACCACGCAGC	
2755	UGGUGAACUU G CGGAA		UUCGG UGAUGGCAUGCACUAUGCGCG AAGUUACACCA	
2770	AGACAGUGGU G AACUU		AAGUU UGAUGGCAUGCACUAUGCGCG ACCACUGUCU	
2780	GAACUUCCU G UAGAA		UUCUA UGAUGGCAUGCACUAUGCGCG AGGGAAGUUC	
2789	UGUAGAAGAC G AGGCC		GGCCU UGAUGGCAUGCACUAUGCGCG GUCUUCUACA	
2813	CACGGCUUUU G UUCAG		CUGAA UGAUGGCAUGCACUAUGCGCG AAAAGCCGUG	
2821	UUUUUCAGAU G CCGGC		GCCGG UGAUGGCAUGCACUAUGCGCG AUCUGAAACA	
2847	UUCUUUUUGU G CGGCC		GGCCG UGAUGGCAUGCACUAUGCGCG ACCAGGGGAA	
2854	GGUGCGGCCU G CUGCU		AGCAG UGAUGGCAUGCACUAUGCGCG AGGCCGCACC	
2857	GGGGCCUGCU G CUGGA		UCCAG UGAUGGCAUGCACUAUGCGCG AGCAGGGCGC	
2881	CCUUGGAGGU G CAGAG		CUCUG UGAUGGCAUGCACUAUGCGCG ACCUCCAGGG	
2888	GGUCAGAGC G ACUAC		GUAGU UGAUGGCAUGCACUAUGCGCG GCUCUGCACC	
2903	CUCACGCUAU G CCCGG		COGGG UGAUGGCAUGCACUAUGCGCG AUAGCUGGAG	
2940	ACCUUCAACC G CCGCU		AGCCG UGAUGGCAUGCACUAUGCGCG GGUUGAAGGU	
2965	GGAGGAACAU G CGUCG		CGACG UGAUGGCAUGCACUAUGCGCG AUGUUCCUCC	
2970	AACAUGCUC G CAAAC		GUUUG UGAUGGCAUGCACUAUGCGCG GACGCAUGUU	
2989	UUGGGGUCUU G CGGCU		AGCCG UGAUGGCAUGCACUAUGCGCG AAGACCCCAA	
2995	UCUUGCGGU G AAGUG		CACUU UGAUGGCAUGCACUAUGCGCG AGCCGCAAGA	
3000	CGGCUGAAGU G UCACA		UGUGA UGAUGGCAUGCACUAUGCGCG ACUUCAGCCG	
3010	GUCACAGCCU G UUUUU		AGAAA UGAUGGCAUGCACUAUGCGCG AGGCUGUGAC	
3022	UUCUGGAUU G CAGGU		ACCUG UGAUGGCAUGCACUAUGCGCG AAUCCAGAA	
3028	AUUUGCAGGU G AACAG		CUGUU UGAUGGCAUGCACUAUGCGCG ACCUGCAAAU	
3046	UCCAGACGGU G UGCAC		GUGCA UGAUGGCAUGCACUAUGCGCG ACCGUCUGGA	
3048	CAGACGGUGU G CACCA		UGGUG UGAUGGCAUGCACUAUGCGCG ACACCGUCUG	
3073	AGAUCCUCCU G CUGCA		UGCAG UGAUGGCAUGCACUAUGCGCG AGGAGGAUCU	
3076	UCCUCCUGCU G CAGGC		GCCUG UGAUGGCAUGCACUAUGCGCG AGCAGGAGGA	
3095	CAGGUUUCAC G CAUGU		ACAUG UGAUGGCAUGCACUAUGCGCG GUGAAACCCUG	
3099	UUUCACGCAU G UGUGC		GCACA UGAUGGCAUGCACUAUGCGCG AUGCGUGAAA	
3101	UCACGCAUGU G UGCUG		CAGCA UGAUGGCAUGCACUAUGCGCG ACAUGCGUGA	
3103	ACGCAUGUGU G CUGCA		UGCAG UGAUGGCAUGCACUAUGCGCG ACACAUGCGU	
3106	CAUGUGUGCU G CAGCU		AGCUG UGAUGGCAUGCACUAUGCGCG AGCACACAUG	

Table 15

3154	CAUUUUUCCU G CCGGU	ACGCG UGAUGGCAUGCACAUAUGCGCG AGGAAAAAUG	
3156	UUUUUCCUGC G CGUCA	UGACG UGAUGGCAUGCACAUAUGCGCG GCAGGAAAAA	
3167	CGUCAUCUCU G ACACG	CGUGU UGAUGGCAUGCACAUAUGCGCG AGAGAUGACG	
3183	GCCUCCUUCU G CUACU	AGUAG UGAUGGCAUGCACAUAUGCGCG AGAGGGAGGC	
3196	ACUCCAUCUCU G AAAGC	GCUUU UGAUGGCAUGCACAUAUGCGCG AGGAUGGAGU	
3209	AGCCAAGAAC G CAGGG	CCCUG UGAUGGCAUGCACAUAUGCGCG GUUCUUGGCU	
3217	ACGCAGGGAU G UCGCU	AGCGA UGAUGGCAUGCACAUAUGCGCG AUCCUUGCGU	
3220	CAGGGAUGUC G CUGGG	CCCAG UGAUGGCAUGCACAUAUGCGCG GACAUCCUUG	
3236	GGCCAAGGCG G CCGCC	GGCGG UGAUGGCAUGCACAUAUGCGCG GCCCUUGGCC	
3239	CAAGGGGGCC G CCGGC	GCCGG UGAUGGCAUGCACAUAUGCGCG GCGGCCCUUG	
3250	CCGGCCUUCU G CCUC	GAGGG UGAUGGCAUGCACAUAUGCGCG AGAGGGCCGG	
3257	UCUGCCUUC G AGGCC	GGCCU UGAUGGCAUGCACAUAUGCGCG GGAGGGCAGA	
3265	CCGAGGCCGU G CAGUG	CACUG UGAUGGCAUGCACAUAUGCGCG ACGGCCUCCG	
3274	UGCAGUGGCU G UGCCA	UGGCA UGAUGGCAUGCACAUAUGCGCG AGCCACUGCA	
3276	CAGUGGCUGU G CCACC	GGUGG UGAUGGCAUGCACAUAUGCGCG ACAGCCACUG	
3292	AAGCAUUCU G CUCAA	UTAGG UGAUGGCAUGCACAUAUGCGCG AGGAAUUCUU	
3301	UGCUCAGCU G ACUCG	CGAGU UGAUGGCAUGCACAUAUGCGCG AGCUUGAGCA	
3306	AAGCUGACUC G ACACC	GGUGU UGAUGGCAUGCACAUAUGCGCG GAGUCAGCUU	
3314	UCGACACCGU G UCACC	GGUGA UGAUGGCAUGCACAUAUGCGCG ACGGUGUGCA	
3325	UCACCUACGU G CCACU	AGUGG UGAUGGCAUGCACAUAUGCGCG ACGUAGGUGA	
3358	CAGCCAGAC G CAGCU	AGCUG UGAUGGCAUGCACAUAUGCGCG GUCUGGGCUG	
3364	AGACGCAGCU G AGUCG	CGACU UGAUGGCAUGCACAUAUGCGCG AGCUGCCUCU	
3385	UCCCGGGGAC G ACGCU	AGCGU UGAUGGCAUGCACAUAUGCGCG GUCCCCGGGA	
3388	CGGGGACGAC G CUGAC	GUCAG UGAUGGCAUGCACAUAUGCGCG GUCGUCUCCG	
3391	GGAGGACGCU G ACUGC	GCAGU UGAUGGCAUGCACAUAUGCGCG AGCGUCGUCC	
3395	GACGUGACU G CCUUG	CAGGG UGAUGGCAUGCACAUAUGCGCG AGUCAGCGUC	
3407	CCUGGAGGCC G CAGCC	GGCUG UGAUGGCAUGCACAUAUGCGCG GGCCUCCAGG	
3424	ACCGGGACU G CCUC	GAGGG UGAUGGCAUGCACAUAUGCGCG AGUGCCGGGU	
3453	AUCCUGGACU G AUGGC	GCCAU UGAUGGCAUGCACAUAUGCGCG AGUCCAGGAU	
3464	AUGGCCACCC G CCAC	GUGGG UGAUGGCAUGCACAUAUGCGCG GGGUGGCCAU	
3479	CAGCCAGGCC G AGAGC	GCUCU UGAUGGCAUGCACAUAUGCGCG GGCCUGGCCUG	

Table 15

3501	CAGCAGCCCU G UCACG	CGUGA UGAUGGCAUGCACUAUGCGCG AGGGCUGCUG	
3506	GCCCUGUAC G CCGGG	CCCAG UGAUGGCAUGCACUAUGCGCG GUGACAGGGC	
3554	ACCCAGGCC G CACCG	CGUG UGAUGGCAUGCACUAUGCGCG GGGCCUGGGU	
3559	GGCCCGACC G CUGGG	CCCAG UGAUGGCAUGCACUAUGCGCG GGUGCGGGCC	
3570	CUGGGAGUCU G AGGCC	GGCCU UGAUGGCAUGCACUAUGCGCG AGACUCCAG	
3577	UCUGAGGCCU G AGUGA	UCACU UGAUGGCAUGCACUAUGCGCG AGGCCUCAGA	
3581	AGGCCUGAGU G AGUGU	ACACU UGAUGGCAUGCACUAUGCGCG ACUCAGGCCU	
3585	CUGAGUGAGU G UUUGG	CCAAA UGAUGGCAUGCACUAUGCGCG ACUCACUCAG	
3593	GUGUUUGGCC G AGGCC	GGCCU UGAUGGCAUGCACUAUGCGCG GGCCAAACAC	
3600	GGCGAGGCCU G CAUGU	ACAUG UGAUGGCAUGCACUAUGCGCG AGGCCUCGGC	
3604	AGGCTUGCAU G UCCGG	CCGGA UGAUGGCAUGCACUAUGCGCG AUGCAGGCCU	
3612	AUGUCCGGCU G AAGGC	GCCUU UGAUGGCAUGCACUAUGCGCG AGCCGGACAU	
3619	GCUGAAGGCU G AGUGU	ACACU UGAUGGCAUGCACUAUGCGCG AGCCUUCAGC	
3623	AAGGCTUGAGU G UCCGG	CCGGA UGAUGGCAUGCACUAUGCGCG ACUCAGCCUU	
3631	GUGUCCGGCU G AGGCC	GGCCU UGAUGGCAUGCACUAUGCGCG AGCCGGACAC	
3638	GCUGAGGCCU G AGCGA	UCGCU UGAUGGCAUGCACUAUGCGCG AGGCCUCAGC	
3642	AGGCTUGAGC G AGUGU	ACACU UGAUGGCAUGCACUAUGCGCG GCUCAGGCCU	
3646	CUAGAGCGAGU G UCCAG	CUGGA UGAUGGCAUGCACUAUGCGCG ACUCGUCACG	
3661	GCCAAAGGCCU G AGUGU	ACACU UGAUGGCAUGCACUAUGCGCG AGCCCUUGGC	
3665	AGGCTUGAGU G UCCAG	CUGGA UGAUGGCAUGCACUAUGCGCG ACUCAGGCCU	
3678	CAGCACACCU G CCGUC	GACGG UGAUGGCAUGCACUAUGCGCG AGGUGUGCUG	
3705	ACAGGCUGGC G CUCGG	CCGAG UGAUGGCAUGCACUAUGCGCG GCCAGCCUUGU	
3789	CCCCAGAUUC G CCAUU	AAUGG UGAUGGCAUGCACUAUGCGCG GAAUCUGGGG	
3795	AUUCGCCAUU G UUCAC	GUGAA UGAUGGCAUGCACUAUGCGCG AAUGGCCAAU	
3806	UUCACCCUC G CCGUG	CAGGG UGAUGGCAUGCACUAUGCGCG GAGGGUGAA	
3811	CCCUUGCCCU G CCGUC	GAGGG UGAUGGCAUGCACUAUGCGCG AGGGCGAGGG	
3821	GGCCUCCUUU G CCGUC	GAAGG UGAUGGCAUGCACUAUGCGCG AAAGGAGGGC	
3854	UGGAGACCCU G AGAAG	CUUCU UGAUGGCAUGCACUAUGCGCG AGGGUCUCCA	
3888	AAUUGGAGU G ACCAA	UUGGU UGAUGGCAUGCACUAUGCGCG ACUCCAAAUU	
3898	GACCAAAGGU G UGCCC	GGGCA UGAUGGCAUGCACUAUGCGCG ACCUUGGUC	
3900	CCAAAGGUGU G CCGUG	CAGGG UGAUGGCAUGCACUAUGCGCG ACACCUUUGG	

Table 15

3905	GGUGUGCCCU G UACAC		GUGUA	UGAUGGCAUGCACUAUGCGCG	AGGGCACACC	
3915	GUACACAGGC G AGGAC		GUCCU	UGAUGGCAUGCACUAUGCGCG	GCCUGUGUAC	
3924	CGAGGACCCU G CACCU		AGGUG	UGAUGGCAUGCACUAUGCGCG	AGGGUCCUCG	
3944	GGGGGUCCCU G UGGGU		ACCCA	UGAUGGCAUGCACUAUGCGCG	AGGGACCCCC	
3966	GGGGGGAGGU G CUGUG		CACAG	UGAUGGCAUGCACUAUGCGCG	ACCUCCCCCC	
3969	GGGAGGUGCU G UGGGA		UCCCA	UGAUGGCAUGCACUAUGCGCG	AGCACCUCCC	
3985	GUAAAAUACU G AAUAU		AUAUU	UGAUGGCAUGCACUAUGCGCG	AGUAUUUUAC	
3993	CUGAAUAUUAU G AGUUU		AAACU	UGAUGGCAUGCACUAUGCGCG	AUAUAUUUCAG	
4008	UUUCAGUUUU G AAAAA		UUUUU	UGAUGGCAUGCACUAUGCGCG	AAAACUGAAA	

Seq1 = TERT (Homo sapiens telomerase reverse transcriptase (TERT) mRNA, 4015 bp); Nakamura *et al.*, Science 277 (5328), 955-959 (1997)
 Input Sequence = TERT. Cut Site = YG/M or UG/U.

Stem Length = 5/10. Core Sequence = UGAUG GCAUGCACUAUGC GCG

Table 16

Table 16: Human telomerase reverse transcriptase (TERT) DNzyme and Target Sequence

nt. Position	DNzyme Sequence	Seq. ID Nos	Substrate	Seq. ID Nos
9	CAGGACGC GGCTAGCTACAACGA AGCGCTGC		GCAGCGCT G GCGTCTCTG	
11	AGCAGGAC GGCTAGCTACAACGA GCAGCGCT		AGCGCTGC G GTCCTGCT	
16	TGGCAGC GGCTAGCTACAACGA AGGACGCA		TGCGTCCT G GCTGCGCA	
19	ACGTGGC GGCTAGCTACAACGA AGCAGGAC		GTCTGCT G GCGCACGT	
21	CCACGTGC GGCTAGCTACAACGA GCAGCAGG		CCTGCTGC G GCACGTGG	
23	TCCACGT GGCTAGCTACAACGA GCGCAGCA		TGCTGGC A ACGTGGGA	
25	CTTCCAC GGCTAGCTACAACGA GTGCGCAG		CTGGCAC G GTGGGAAG	
32	GCCAGGCG GGCTAGCTACAACGA TTCCACAG		CGTGGAA G GCCCTGGC	
38	GCCGGGCG GGCTAGCTACAACGA CAGGGCTT		AAGCCCTG G GCCCGGCG	
44	GCGGTGGC GGCTAGCTACAACGA CGGGGCCA		TGGCCCGG G GCCACCCG	
47	GCGGGGT GGCTAGCTACAACGA GCGCGGG		CCCCGGCC A ACCCCCGC	
53	GGCATCGC GGCTAGCTACAACGA GCGGTGG		CCACCCCG G GCGATGCG	
56	CGGGGCAT GGCTAGCTACAACGA CCGGGGG		CCCCCGCG A ATGCCGCG	
58	CGCGCGC GGCTAGCTACAACGA ATCGGGG		CCCGCGAT G GCCGCGCG	
61	GAGCGCGC GGCTAGCTACAACGA GGCATCGC		GCGATGCC G GCGGCTC	
63	GGGAGCGC GGCTAGCTACAACGA GCGGCATC		GATGCCG G GCGTCCC	
65	CGGGGAGC GGCTAGCTACAACGA GCGCGGCA		TGCCGCG G GCTCCCG	
72	TCGGCAGC GGCTAGCTACAACGA GGGGAGCG		CGTCCCC G GCTGCCGA	
75	GGCTCGC GGCTAGCTACAACGA AGCGGGGA		TCCCCGCT G GCGAGGCC	
80	CGCACGGC GGCTAGCTACAACGA TCGGCAGC		GCTGCCGA G GCGGTGG	
83	GAGCGCAC GGCTAGCTACAACGA GGTCCGGC		GCCGAGCC G GTGCGCTC	
85	GGGAGCGC GGCTAGCTACAACGA ACGGCTCG		CGAGCGT G GCGTCCC	
87	CAGGGAGC GGCTAGCTACAACGA GCACGGCT		AGCCGTG G GCTCCCTG	
94	TGCGCAGC GGCTAGCTACAACGA AGGAGCG		CGTCCCT G GCTGGCA	
97	GCGTGGC GGCTAGCTACAACGA AGCAGGGA		TCCCTGCT G GCGCAGCC	
99	GTGGCTGC GGCTAGCTACAACGA GCAGCAGG		CCTGCTG G GCAGCCAC	
102	GTAGTGGC GGCTAGCTACAACGA TGCGCAGC		GCTGCCGA G GCCACTAC	

Table 16

105	GCGTAGT	GGTAGCTACAACGA	GGTGCGC	GCGCAGC A	ACTACGCG	
108	CTCGGGT	GGTAGCTACAACGA	AGTGCTG	CAGCCACT A	ACGCGGAG	
111	CACCTGC	GGTAGCTACAACGA	GGTAGTG	CCACTACC G	GCGAGGTG	
116	GGCAGCAC	GGTAGCTACAACGA	CTCGGGT	ACCGCGAG G	GTGCTGCC	
118	GCGGCAGC	GGTAGCTACAACGA	ACCTGCG	GCGAGGT G	GCTGCCGC	
121	CCAGCGGC	GGTAGCTACAACGA	AGCACTC	GAGGTGCT G	GCGCTGG	
124	TGGCAGC	GGTAGCTACAACGA	GGCAGCAC	GTGCTGCC G	GCTGGCCA	
128	AACGTGGC	GGTAGCTACAACGA	CAGCGCA	TGCCGCTG G	GCCACGTT	
131	ACGAAGT	GGTAGCTACAACGA	GGCCAGC	CGCTGGCC A	ACGTTCTG	
133	GCACGAAC	GGTAGCTACAACGA	GTGGCCAG	CTGGCCAC G	GTTCTGTG	
137	GCGGCAC	GGTAGCTACAACGA	GAACGTG	CCACGTTC G	GTGCGGCG	
139	GGCGCGC	GGTAGCTACAACGA	ACGAAGT	ACGTTCTG G	GCGGCGCC	
142	CCAGCGC	GGTAGCTACAACGA	CGCAGAA	TTCGTGCG G	GCGCTGG	
144	CCCCAGC	GGTAGCTACAACGA	GCCGACG	CGTCGCG G	GCTGGGG	
151	CCTGGGC	GGTAGCTACAACGA	CCCAGCG	CGCTGGG G	GCCCCAGG	
159	CCGCCAGC	GGTAGCTACAACGA	CCTGGGC	GCCCCAGG G	GCTGGCGG	
163	CCAGCGC	GGTAGCTACAACGA	CAGCCCTG	CAGGGCTG G	GCGCTGG	
166	GCACCAGC	GGTAGCTACAACGA	GCCAGCC	GCTGGCG G	GCTGTTGC	
170	CGCTGCAC	GGTAGCTACAACGA	CAGCGCC	GCGGGCTG G	GTGCAGCG	
172	GCGCTGC	GGTAGCTACAACGA	ACCAGCG	CGCTGGT G	GCAGCGCG	
175	CCCCGCGC	GGTAGCTACAACGA	TGCACAG	CTGGTGCA G	GCGCGGG	
177	GTCCCGC	GGTAGCTACAACGA	GCTGCACC	GGTGCAGC G	GCGGGGAC	
183	GCGCGGT	GGTAGCTACAACGA	CCCCGCG	GCGCGGG A	ACCCGGCG	
188	AAAGCCG	GGTAGCTACAACGA	CGGGTCCC	GGACCCG G	GCGCTTT	
191	CGGAAAGC	GGTAGCTACAACGA	CGCCGGT	ACCCGGCG G	GCTTCCG	
198	CAGCGGC	GGTAGCTACAACGA	GGAAAGC	GGCTTCC G	GCGGCTG	
200	ACCAGCGC	GGTAGCTACAACGA	GCGAAAG	CTTCCGCG G	GCGTGGT	
202	CCACCAGC	GGTAGCTACAACGA	GCGGGAA	TTCGCGCG G	GCTGGTGG	
206	TGGGCCAC	GGTAGCTACAACGA	CAGCGCG	GCGGCTG G	GTGGCCCA	
209	CACTGGGC	GGTAGCTACAACGA	CACCAAG	CGCTGGT G	GCCAGTGG	
214	CCAGGCAC	GGTAGCTACAACGA	TGGCCAC	GTGGCCCA G	GTGCTGG	

Table 16

216	CACCAGGC	GGCTAGCTACAACGA	ACTGGGCC		GGCCAGT G	GCCTGGTG	
221	ACGCACAC	GGCTAGCTACAACGA	CAGGCACT		AGTGCTG G	GTGTGCGT	
223	GCACGCAC	GGCTAGCTACAACGA	ACCAGGCA		TGCCTGGT G	GTGCGTGC	
225	GGCACGCG	GGCTAGCTACAACGA	ACACCAGG		CCTGTGTG	G GGTGCC	
227	CAGGGCAC	GGCTAGCTACAACGA	GCACACCA		TGGTGTGC G	GTGCCCTG	
229	CCCAGGGC	GGCTAGCTACAACGA	ACGCACAC		GTGTGCGT G	GCCCTGGG	
237	CGGTGCGT	GGCTAGCTACAACGA	CCCAGGGC		GCCCTGGG A	ACGCACCG	
239	GGCGGTGC	GGCTAGCTACAACGA	GTCCCAGG		CCTGGGAC G	GCACGGGC	
241	GGGCCCGT	GGCTAGCTACAACGA	CGTCCCA		TGGGACGC A	ACGGCCGC	
244	GGGGCGGC	GGCTAGCTACAACGA	CGTCCGTC		GACGCACG G	GCCGCCCC	
247	CGGGGGGC	GGCTAGCTACAACGA	GGCGGTGC		GCACGGCC G	GCCCCCGG	
254	GGGGCGGC	GGCTAGCTACAACGA	GGGGGGCG		GCCCCCCC G	GCCCCCCC	
257	GAGGGGGC	GGCTAGCTACAACGA	GGCGGGGG		CCCCCGCC G	GCCCCCTC	
270	CACCTGGC	GGCTAGCTACAACGA	GGAAGGAG		CTCCTTCC G	GCCAGGTG	
275	CAGGACAC	GGCTAGCTACAACGA	CTGGCGGA		TCCGCCAG G	GTGTCTCTG	
277	GGCAGGAC	GGCTAGCTACAACGA	ACCTGGCG		CGCCAGGT G	GTCTTGCC	
282	CTTCAGGC	GGCTAGCTACAACGA	AGGACACC		GGTGTCTT G	GCCTGAAG	
292	CCACCAGC	GGCTAGCTACAACGA	TCCTTCAG		CTGAAGGA G	GCTGGTGG	
296	CGGGCCAC	GGCTAGCTACAACGA	CAGTCTCT		AGGAGCTG G	GTGGCCCG	
299	ACTCGGGC	GGCTAGCTACAACGA	CACCAGCT		AGTGTGTG G	GCCCGAGT	
305	TGCAGCAC	GGCTAGCTACAACGA	TCGGGCCA		TGGCCCCA G	GTGCTGCA	
307	TCTGCAGC	GGCTAGCTACAACGA	ACTCGGGC		GCCCGAGT G	GCTGCAGA	
310	GCCTCTGC	GGCTAGCTACAACGA	AGCACTCG		CGAGTGTCT G	GCAGAGGC	
316	CGCACAGC	GGCTAGCTACAACGA	CTCTGCAG		CTGCAGAG G	GCTGTGCG	
319	GCTCGCAC	GGCTAGCTACAACGA	AGCTCTGT		CAGAGGCT G	GTSCGAGC	
321	GCCTCGC	GGCTAGCTACAACGA	ACAGCTCT		GAGGCTGT G	GCGAGCGC	
325	CGCGGGCG	GGCTAGCTACAACGA	TCGCACAG		CTGTGCGA G	GCGCGGGC	
327	CGCGCCCG	GGCTAGCTACAACGA	GCTCGCAC		GTGCGAGC G	GCGCGCGG	
330	CTTCGCGC	GGCTAGCTACAACGA	CGCGCTCG		CGAGCGCG G	GCGCGAAG	
332	TTCTTCGC	GGCTAGCTACAACGA	GCCGCGCT		AGCGCGCG G	GCGAAGAA	
339	CAGCACGT	GGCTAGCTACAACGA	TCTTCGCG		CGCGAAGA A	ACGTGCTG	

Table 16

341	GCCAGCAC	GGCTAGCTACAACGA	GTTCTTGG		CGAAGAAC	G	GTGCTGGC	
343	AGGCCAGC	GGCTAGCTACAACGA	ACGTTCTT		AAGAACGT	G	GCTGGCCT	
347	CCGAAGGC	GGCTAGCTACAACGA	CAGCACGT		ACGTGCTG	G	GCCTTCGG	
354	CGCGAAGC	GGCTAGCTACAACGA	CGAAGGCC		GGCCTTCG	G	GCTTCGGG	
359	AGCAGCGC	GGCTAGCTACAACGA	GAAGCCGA		TCGGCTTC	G	GGCTGCT	
361	CCAGCAGC	GGCTAGCTACAACGA	GCGAAGCC		GGCTTCGC	G	GCTGCTGG	
364	CGTCCAGC	GGCTAGCTACAACGA	AGCGCGAA		TTGCGCGT	G	GCTGGACG	
369	GGCCCCGT	GGCTAGCTACAACGA	CCAGCAGC		GCTGCTGG	A	ACGGGGCC	
374	CCGCGGGC	GGCTAGCTACAACGA	CCGCTCCA		TGGACGGG	G	GCCCCGGG	
378	GCCCCCGC	GGCTAGCTACAACGA	GGGCCCGG		CGGGGGCC	G	GGGGGGGC	
384	GGGGGGGC	GGCTAGCTACAACGA	CCCGCGGG		CCGCGGGG	G	GGCCCCCC	
395	GTGAAGGC	GGCTAGCTACAACGA	CTCGGGGG		CCCCCGAG	G	GCCTTCAC	
401	CTGCTGGT	GGCTAGCTACAACGA	GAAGGCCT		AGGCCTTC	A	ACCACCAG	
404	ACGCTGGT	GGCTAGCTACAACGA	GGTGAAGG		CCTTCACC	A	ACCAGCGT	
408	GCGCACGC	GGCTAGCTACAACGA	TGTTGGTG		CACCACCA	G	GCGTGCGC	
410	CTGCGCAC	GGCTAGCTACAACGA	GCTGTTGG		CCACCAGC	G	GTGCGCAG	
412	AGTGGCGC	GGCTAGCTACAACGA	ACGTTGGT		ACCAGCGT	G	GCGCAGCT	
414	GTAGCTGC	GGCTAGCTACAACGA	GCAGGCTG		CAGCGTGC	G	GCAGCTAC	
417	CAGGTAGC	GGCTAGCTACAACGA	TGCGCACG		CGTGCGCA	G	GCTACCTG	
420	GGGCAGGT	GGCTAGCTACAACGA	AGCTGCGC		GCGCAGCT	A	ACCTGCCC	
424	TGTTGGGC	GGCTAGCTACAACGA	AGGTAGCT		AGCTACCT	G	GGCCCAACA	
429	CACCGTGT	GGCTAGCTACAACGA	TGGGCAGG		CCTGCCCA	A	ACACGGTG	
431	GTCAACCGT	GGCTAGCTACAACGA	GTTGGGCA		TGCCCAAC	A	ACGGTGAC	
434	TCGGTCAC	GGCTAGCTACAACGA	CGTGTGGG		CCAACACG	G	GTGACCGA	
437	GCGTCGGT	GGCTAGCTACAACGA	CACGTTGT		ACACGGTG	A	ACCGACGC	
441	CAGTGGCT	GGCTAGCTACAACGA	CGGTCACC		GGTGACCG	A	ACGCACTG	
443	CGCAGTGC	GGCTAGCTACAACGA	GTCGGTCA		TGACCCAG	G	GCACTGCG	
445	CCCGCAGT	GGCTAGCTACAACGA	GCGTCGGT		ACCGACGC	A	ACTGCGGG	
448	TCCCCCGC	GGCTAGCTACAACGA	AGTGCCTC		GACGCACT	G	GCGGGGGA	
456	CGCCCCCGC	GGCTAGCTACAACGA	TCCCCCCGC		GCGGGGGA	G	GCGGGGCG	
461	CCCCACGC	GGCTAGCTACAACGA	CCCGCTCC		GGAGCGGG	G	GCGTGGGG	

Table 16

463	GCCCCAC	GGTAGCTACAACGA	GCCCCGCT		AGCGGGC G GTGGGGC
469	GCAGAGC	GGTAGCTACAACGA	CCCCACG		GCGTGGG G GCTGCTG
472	GCAGAGC	GGTAGCTACAACGA	AGCCCCA		TGGGGGCT G GCTGCTG
475	GGCGCAGC	GGTAGCTACAACGA	AGAGCCC		GCGCTGCT G GCTGCGC
478	CGCGGCGC	GGTAGCTACAACGA	AGCAGAG		CTGCTGCT G GCGCCGCG
480	CACGCGGC	GGTAGCTACAACGA	GCAGCAGC		GCTGCTGC G GCGCGCTG
483	GCCACGCG	GGTAGCTACAACGA	GGCGCAGC		GCTGGCC G GCGTGGC
485	TCGCCAC	GGTAGCTACAACGA	GCGGCGCA		TGGCCGC G GTGGCGA
489	GTCGTCG	GGTAGCTACAACGA	CCACGCG		CCCGTGG G GCGAGGAC
492	CAGTCGT	GGTAGCTACAACGA	CGCCACG		CGTGGCG A ACGAGTG
495	CAGCAGT	GGTAGCTACAACGA	CGTCGCC		GCGCGAG A ACGTCTG
497	ACCAGCAC	GGTAGCTACAACGA	GTCGTGC		GCGACGAC G GTGCTGT
499	GAACGAGC	GGTAGCTACAACGA	ACGTCTC		GACGACGT G GCTGTTT
503	AGGTGAAC	GGTAGCTACAACGA	CAGCAGT		ACGTGCTG G GTTACCT
507	CAGCAGGT	GGTAGCTACAACGA	GAACGAGC		GCTGGTC A ACGTCTG
511	GTGCCAGC	GGTAGCTACAACGA	AGGTGAAC		GTTACCT G GCTGGAC
515	CAGCGTC	GGTAGCTACAACGA	CAGCAGT		ACGTGCTG G GCACGCTG
517	CGCAGCGT	GGTAGCTACAACGA	GCCAGCAG		CTGCTGGC A ACGTGGG
519	CGCGCAGC	GGTAGCTACAACGA	GTGCCAGC		GCTGGCAC G GCTGCGG
522	GAGCGCGC	GGTAGCTACAACGA	AGCGTGC		GGCAGGCT G GCGGCTC
524	AAGAGCGC	GGTAGCTACAACGA	GCAGCGTG		CACGCTGC G GCGCTCT
526	CAAGAGGC	GGTAGCTACAACGA	GCGCAGCG		CGCTGGC G GCTCTTTG
533	ACCAGCAC	GGTAGCTACAACGA	AAAGAGCG		CGCTCTTT G GTGCTGGT
535	CCACCAGC	GGTAGCTACAACGA	ACAAAGAG		CTCTTTGT G GCTGGTG
539	GGAGCCAC	GGTAGCTACAACGA	CAGCACAA		TTGTGCTG G GTGGCTC
542	CTGGGAGC	GGTAGCTACAACGA	CACCAGCA		TGCTGGTG G GCTCCAG
549	GGCGCAGC	GGTAGCTACAACGA	TGGAGCGC		GGTCCCA G GCTGGCC
552	GTAGGCGC	GGTAGCTACAACGA	AGCTGGGA		TCCAGCT G GCGCTAC
554	TGGTAGGC	GGTAGCTACAACGA	GCAGCTGG		CCAGCTGC G GCTTACCA
558	CACCTGGT	GGTAGCTACAACGA	AGGCGCAG		CTGGGCT A ACCAGGTG
563	CCGACAC	GGTAGCTACAACGA	CTGTAGG		CCTACCAG G GTGTGGG

Table 16

565	GCCCGCAC	GGCTAGCTACAACGA	ACCTGGTA		TACCAGGT G	GTGCGGCG
567	CGGCCCGC	GGCTAGCTACAACGA	ACACCTGG		CCAGGTGT G	GCGGGCGG
571	GCGGCGGC	GGCTAGCTACAACGA	CCGCACAC		GTGTGGG G	GCGGCGGC
574	ACAGCGGC	GGCTAGCTACAACGA	GGCCCGCA		TGCGGGCC G	GCGGCTGT
577	GGTACAGC	GGCTAGCTACAACGA	GCGGGCCC		GGGCGGCC G	GCTGTACC
580	GCTGGTAC	GGCTAGCTACAACGA	AGCGGCGG		CCGCCGCT G	GTACCAGC
582	GAGCTGGT	GGCTAGCTACAACGA	ACAGCGGC		GCCGCTGT A	ACCAGCTC
586	CGCCGAGC	GGCTAGCTACAACGA	TGCTACAG		CTGTACCA G	GCTCGGCG
591	GGCAGCGC	GGCTAGCTACAACGA	CGAGCTGG		CCAGCTCG G	GCGCTGCC
593	GTGGCAGC	GGCTAGCTACAACGA	GCGGAGCT		AGCTCGGC G	GCTGCCAC
596	TGAGTGGC	GGCTAGCTACAACGA	AGCGCCGA		TCGGGCGT G	GCCACTCA
599	GCTTGAGT	GGCTAGCTACAACGA	GGCAGCGC		GCGCTGCC A	ACTCAGGC
605	GCCCGGGC	GGCTAGCTACAACGA	CTGAGTGG		CCACTCAG G	GCCCGGCC
610	GCGGGGGC	GGCTAGCTACAACGA	CGGGCCTG		CAGGCCCG G	GCCCCCGC
616	CGTGTGGC	GGCTAGCTACAACGA	GGGGGCCG		CGGCCCCC G	GCCACACG
619	TAGCGTGT	GGCTAGCTACAACGA	GCGGGGGG		CCCCCGCC A	ACACGCTA
621	ACTAGCGT	GGCTAGCTACAACGA	GTGGCGGG		CCCGCCAC A	ACGCTAGT
623	CCACTAGC	GGCTAGCTACAACGA	GTGTGGCG		CGCCACAC G	GCTAGTGG
627	GGGTCCAC	GGCTAGCTACAACGA	TAGCGTGT		ACACGCTA G	GTGGACCC
631	TTCGGGGT	GGCTAGCTACAACGA	CCACTAGC		GCTAGTGG A	ACCCCGAA
640	CCAGACGC	GGCTAGCTACAACGA	CTTCGGGG		CCCCGAAG G	GCGTCTGG
642	TCCCAGAC	GGCTAGCTACAACGA	GCTTCGG		CCGAAGGC G	GTCTGGGA
649	GTTCGCAT	GGCTAGCTACAACGA	CCAGACGC		CGTCTGGG A	ATCGGAAC
651	CCGTTCCG	GGCTAGCTACAACGA	ATCCAGA		TCTGGGAT G	GCGAACGG
655	AGGCCCGT	GGCTAGCTACAACGA	TCCATCC		GGATGCGA A	ACGGGCCT
659	TTCCAGGC	GGCTAGCTACAACGA	CCGTTCCG		GCGAACGG G	GCCTGGAA
666	GCTATGGT	GGCTAGCTACAACGA	TCCAGGCC		GGCCTGGA A	ACCATAGC
669	GACGCTAT	GGCTAGCTACAACGA	GGTTCACG		CTGGAACC A	ATAGCGTC
672	CCTGACGC	GGCTAGCTACAACGA	TATGGTTC		GAACCAT A	GCGTCAGG
674	TCCTTGAC	GGCTAGCTACAACGA	GCTATGGT		ACCATAGC G	GTACAGGA
683	ACCCCGGC	GGCTAGCTACAACGA	CTCCCTGA		TCAGGGAG G	GCCGGGGT

Table 16

689	AGGGGAC	GGTAGCTACAACGA	CCCGGCTT		AGCCCGG G GTCCCCCT	
699	TGCAGGC	GGTAGCTACAACGA	CCAGGGG		CCCCCTGG G GCCTGCCA	
703	GGCTGGC	GGTAGCTACAACGA	AGCCCCAG		CTGGGCTT G GCCAGCCC	
707	CCCGGGC	GGTAGCTACAACGA	TGCAGGC		GCCTGCCA G GCCCCGGG	
714	CCTCGAC	GGTAGCTACAACGA	CCGGGGCT		AGCCCGG G GTGCGAGG	
716	CTCCTGC	GGTAGCTACAACGA	ACCCGGG		CCCGGGT G GCGAGGAG	
724	CCCGGCG	GGTAGCTACAACGA	CTCCTGC		GCGAGGAG G GCGGGG	
726	GCCCCCG	GGTAGCTACAACGA	GCCTCCTC		GAGGAGG G GCGGGGC	
732	GGCACTGC	GGTAGCTACAACGA	CCCGGCG		GCGCGGG G GCAGTGCC	
735	GCTGGCAC	GGTAGCTACAACGA	TGCCCCCG		CGGGGGA G GTGCCAGC	
737	CGGCTGGC	GGTAGCTACAACGA	ACTGCCCC		GGGGCAGT G GCCAGCCG	
741	ACTTCGGC	GGTAGCTACAACGA	TGGCACTG		CAGTGCCA G GCCAAGT	
747	CGGCAGAC	GGTAGCTACAACGA	TTGGGCTG		CAGCCGAA G GTCTGCCG	
751	GCAACGGC	GGTAGCTACAACGA	AGACTTCG		CGAAGTCT G GCGTTGC	
754	TGGGCAAC	GGTAGCTACAACGA	GGCAGACT		AGTCTGCC G GTTGCCCA	
757	TCTTGGGC	GGTAGCTACAACGA	AACGGCAG		CTGCCGTT G GCCAAGA	
766	GCCTGGGC	GGTAGCTACAACGA	CTCTTGGG		CCCAAGAG G GCCAGGC	
772	CGCCACGC	GGTAGCTACAACGA	CTGGGCTT		AGGCCAG G GCGTGGCG	
774	AGCGCCAC	GGTAGCTACAACGA	GCCTGGGC		GCCCAGC G GTGGCGCT	
777	GGCAGCGC	GGTAGCTACAACGA	CACGCTTG		CAGGCGTG G GCGTGCC	
779	GGGCAGC	GGTAGCTACAACGA	GCCAGGCC		GGGCTGGC G GCTGCCCC	
782	TCAGGGGC	GGTAGCTACAACGA	AGCGCCAC		GTGGCGCT G GCCCTGA	
790	GCTCCGGC	GGTAGCTACAACGA	TCAGGGGC		GCCCCGA G GCCGAGC	
796	GCGTCGCG	GGTAGCTACAACGA	TCCGGCTC		GAGCCGA G GCGAGCGC	
800	ACGGCGT	GGTAGCTACAACGA	CCGCTCCG		CGGAGCGG A ACGCCCGT	
802	CAACGGGC	GGTAGCTACAACGA	GTCCGCTC		GAGCGGAC G GCCGTTG	
806	TGCCCAAC	GGTAGCTACAACGA	GGGGTCC		GGACGCC G GTTGGGA	
811	ACCCCTGC	GGTAGCTACAACGA	CCAAGGG		CCGTTGG G GCAGGGT	
817	CCCAGGAC	GGTAGCTACAACGA	CCCTGCC		GGCAGGG G GTCTGGG	
824	GGGTGGGC	GGTAGCTACAACGA	CCAGGACC		GGTCTGG G GCCACCC	
828	GCCCGGGT	GGTAGCTACAACGA	GGGCCAG		CTGGGCC A ACCGGGC	

Table 16

834	CGTCTGC	GGCTAGCTACAACGA	CCGGTGG		CCACCCG G GCAGACG
839	CCACGCGT	GGCTAGCTACAACGA	CCTGCCG		CGGCAGG A ACGGTGG
841	GTCCACGC	GGCTAGCTACAACGA	GTCTGCC		GGCAGGAC G GCGTGGAC
843	CGGTCCAC	GGCTAGCTACAACGA	CGTCTCTG		CAGGACGC G GTGGACCG
847	CACTCGGT	GGCTAGCTACAACGA	CCACGGT		ACGCGTGG A ACCGAGTG
852	ACGGTCCAC	GGCTAGCTACAACGA	TCGGTCCA		TGGACCGA G GTGACCGT
855	ACCACGGT	GGCTAGCTACAACGA	CACTCGGT		ACCGAGTG A ACCGTGGT
858	GAACCCAC	GGCTAGCTACAACGA	GGTCACTC		GAGTGACC G GTGGTTTC
861	ACAGAAAC	GGCTAGCTACAACGA	CACGGTCA		TGACCGTG G GTTCTGT
867	CACCACAC	GGCTAGCTACAACGA	AGAAACCA		TGGTTTCT G GTGTGGTG
869	GACACCCAC	GGCTAGCTACAACGA	ACAGAAAC		GTCTCTGT G GTGTGTC
872	GGTGACAC	GGCTAGCTACAACGA	CACACAGA		TCTGTGTG G GTGTACCC
874	CAGGTGAC	GGCTAGCTACAACGA	ACCACACA		TGTGTGGT G GTCACTG
877	TGGCAGGT	GGCTAGCTACAACGA	GACACCAC		GTGGTGTG A ACCTGCCA
881	GGTCTGGC	GGCTAGCTACAACGA	AGGTGACA		TGTCACCT G GCCAGACC
886	CGCGGGT	GGCTAGCTACAACGA	CTGGCAGG		CCTGCCAG A ACCGCGG
890	TCTTCGGC	GGCTAGCTACAACGA	GGGTCTGG		CCAGACCC G GCCGAGA
899	GAGGTGGC	GGCTAGCTACAACGA	TTCTTCGG		CCGAGAA G GCCACCTC
902	AAAGAGGT	GGCTAGCTACAACGA	GGCTTCTT		AAGAAGCC A ACCTCTTT
915	GAGCGCAC	GGCTAGCTACAACGA	CCTCCAAA		TTTGGAGG G GTGCGTC
917	GAGAGCGC	GGCTAGCTACAACGA	ACCCTCCA		TGGAGGGT G GCGTCTC
919	CAGAGAGC	GGCTAGCTACAACGA	GCACCCCTC		GAGGTGC G GCTCTCTG
927	CGCGGTGC	GGCTAGCTACAACGA	CAGAGAGC		GCTCTCTG G GCACGCGC
929	TGGCGCGT	GGCTAGCTACAACGA	GCCAGAGA		TCTCTGGC A ACGCGCCA
931	AGTGGCGC	GGCTAGCTACAACGA	GTGCCAGA		TCTGGCAC G GCGCCACT
933	GGAGTGGC	GGCTAGCTACAACGA	GGGTGCCA		TGGCACGC G GCCACTCC
936	GTGGGAGT	GGCTAGCTACAACGA	GGCGGTG		CACGGCC A ACTCCAC
942	GGATGGGT	GGCTAGCTACAACGA	GGGAGTGG		CCACTCCC A ACCATCC
946	CCACGGAT	GGCTAGCTACAACGA	GGGTGGGA		TCCACCCC A ATCGTGG
950	CGGCCCCAC	GGCTAGCTACAACGA	GGATGGGT		ACCATCC G GTGGGCCG
954	CTGGCGGC	GGCTAGCTACAACGA	CCACGGAT		ATCGTGG G GCCGCCAG

Table 16

957	GTGCTGGC	GGCTAGCTACAACGA	GGCCACG		CGTGGGC G	GCCAGCAC	
961	CGTGTGC	GGCTAGCTACAACGA	TGGCGGC		GGCGCCA G	GCACCAAG	
963	CGCGTGGT	GGCTAGCTACAACGA	GCTGGCGG		CGCCAGC A	ACCAAGCG	
966	GCCCGCGT	GGCTAGCTACAACGA	GGTGTGG		CCAGACC A	ACGGGGC	
968	GGGCGCGC	GGCTAGCTACAACGA	GTGTGCT		AGCACAC G	GCGGGCCC	
972	TGGGGGC	GGCTAGCTACAACGA	CCGCTGG		CCAGCGG G	GCCCCCA	
979	ATGTGGAT	GGCTAGCTACAACGA	GGGGGCC		GGCCCCC A	ATCCACAT	
983	CGCGATGT	GGCTAGCTACAACGA	GGATGGG		CCCCATC A	ACATCGCG	
985	GCCGGAT	GGCTAGCTACAACGA	GTGGATG		CCATCCAC A	ATCGGGC	
988	GTGGCCGC	GGCTAGCTACAACGA	GATGTGA		TCCACATC G	GCGGCCAC	
991	GTGTGGC	GGCTAGCTACAACGA	CGCATGT		ACATCGCG G	GCACCCAC	
994	GACGTGGT	GGCTAGCTACAACGA	GGCCGCGA		TCGGGGC A	ACCAAGTC	
997	AGGACGT	GGCTAGCTACAACGA	GGTGCCG		CGGCCAC A	AGTCCCT	
999	CCAGGGAC	GGCTAGCTACAACGA	GTGTGGC		GCCACCAC G	GTCCCTGG	
1008	AGCGTGT	GGCTAGCTACAACGA	CCCAGGA		TCCCTGG A	ACACGCCT	
1010	CAAGCGT	GGCTAGCTACAACGA	GTCCAGG		CCTGGGAC A	ACGCCTTG	
1012	GACAAGC	GGCTAGCTACAACGA	GTGTCCA		TGGGACAC G	GCCTTGTC	
1017	CGGGGAC	GGCTAGCTACAACGA	AAGCGTG		CAGCCIT G	GTCCCCCG	
1025	GCGTACAC	GGCTAGCTACAACGA	CGGGGAC		GTCCCCG G	GTGTAGC	
1027	CGCGTAC	GGCTAGCTACAACGA	ACCGGGG		CCCCCGT G	GTAGGCCG	
1029	CTCGCGT	GGCTAGCTACAACGA	ACACGGG		CCGGTGT A	ACGCCAG	
1031	GTCTCGC	GGCTAGCTACAACGA	GTACACG		CGGTGTAC G	GCCGAGAC	
1037	TGCTTGGT	GGCTAGCTACAACGA	CTCGCGT		ACGCCGAG A	ACCAAGCA	
1042	GGAAGTC	GGCTAGCTACAACGA	TTGTCTC		GAGACCAA G	GCATTTC	
1044	GAGGAAT	GGCTAGCTACAACGA	GCTTGGTC		GACCAAGC A	ACTTCCTC	
1053	TGAGAGT	GGCTAGCTACAACGA	AGAGGAAG		CTTCTCT A	ACTCCTCA	
1062	CTTGTGCG	GGCTAGCTACAACGA	CTAGGAG		CTCCTCAG G	GCGACAAG	
1065	CTCCTTGT	GGCTAGCTACAACGA	CGCTGAG		CTCAGGG A	ACAAGGAG	
1072	GCAGCTGC	GGCTAGCTACAACGA	TCCTTGT		GACAAAG A	GCAAGTGC	
1075	GCCGCAGC	GGCTAGCTACAACGA	TGCTCCT		AAGGAGCA G	GCTGCGGC	
1078	AGGGCCGC	GGCTAGCTACAACGA	AGTGTCT		GAGCAGCT G	GCGGCCCT	

Table 16

1081	AGGAGGGC GGCTAGCTACAACGA CGCAGCTG		CAGCTGGC G GCCCTCCT
1093	AGCTGAGT GGCTAGCTACAACGA AGGAAGGA		TCCTTCTT A ACTCAGCT
1098	CAGAGAGC GGCTAGCTACAACGA TGAGTAGG		CCTACTCA G GCTCTCTG
1108	GGCTGGGC GGCTAGCTACAACGA CTCAGAGA		TCTCTGAG G GCCCAGCC
1113	AGTCAGGC GGCTAGCTACAACGA TGGGCCTC		GAGGCCCA G GCCTGACT
1118	GCGCCAGT GGCTAGCTACAACGA CAGCTGG		CCAGCCTG A ACTGGCGC
1122	CCGAGCGC GGCTAGCTACAACGA CAGTCAGG		CCTGACTG G GCGCTCGG
1124	CTCCGAGC GGCTAGCTACAACGA GCCAGTCA		TGACTGGC G GCTCGGAG
1132	CCACGAGC GGCTAGCTACAACGA CTCGAGC		GCTCGGAG G GCTCGTGG
1136	GTCTCCAC GGCTAGCTACAACGA GAGCTCC		GGAGGCTC G GTGGAGAC
1142	AAGATGGT GGCTAGCTACAACGA CTCACAGA		TCGTGGAG A ACCATCTT
1145	AGAAAGAT GGCTAGCTACAACGA GGTCTCCA		TGGAGACC A ATCTTTCT
1155	CCTGGAAC GGCTAGCTACAACGA CCAGAAAG		CTTTCTGG G GTTCCAGG
1162	TCCAGGGC GGCTAGCTACAACGA CTGGAACC		GGTTCCAG G GCCTGGA
1169	CCTGGCAT GGCTAGCTACAACGA CCAGGGCC		GGCCCTGG A ATGCCAGG
1171	TCCCTGGC GGCTAGCTACAACGA ATCCAGGG		CCCTGGAT G GCCAGGA
1178	CGGGGAGT GGCTAGCTACAACGA CCCTGGCA		TGCCAGGG A ACTCCCCG
1185	CAACTGTC GGCTAGCTACAACGA GGGGAGTC		GACTCCCC G GCAGGTTG
1189	GGGGCAAC GGCTAGCTACAACGA CTGGGGG		CCCCGCAG G GTTGCCCC
1192	GGCGGGC GGCTAGCTACAACGA AACCTGCG		CGCAGGTT G GCCTGCCC
1197	GGGCAGGC GGCTAGCTACAACGA GGGGCAAC		GTTCGCCC G GCCTGCCC
1201	GCTGGGGC GGCTAGCTACAACGA AGGCGGGG		CCCGGCTT G GCCTGCCC
1207	AGTAGGCG GGCTAGCTACAACGA TGGGGCAG		CTGGCCCC G GCGTACT
1209	CCAGTAGC GGCTAGCTACAACGA GCTGGGGC		GGCCGAGC G GCTACTGG
1212	TTGCCAGT GGCTAGCTACAACGA AGCGCTGG		CCAGCGCT A ACTGGCAA
1216	GCAATTGC GGCTAGCTACAACGA CAGTAGCG		CGCTACTG G GCAAATGC
1220	GGCCGCAT GGCTAGCTACAACGA TTGCCAGT		ACTGGCAA A ATGCGGCC
1222	GGGGCGGC GGCTAGCTACAACGA ATTTGCCA		TGGCAAAAT G GCGGCCCC
1225	ACAGGGGC GGCTAGCTACAACGA CGCATTTG		CAAAATGCG G GCCCTGT
1231	CCAGAAAC GGCTAGCTACAACGA AGGGCGCG		CGGCCCTT G GTTCTCTG
1240	CAAGCAGC GGCTAGCTACAACGA TCCAGAAA		TTTCTGGA G GCTGCTTG

Table 16

1243	TCCCAAGC	GGCTAGCTACAACGA	AGCTCCAG		CTGGAGCT G	GCTTGGGA	
1251	CGCGTGGT	GGCTAGCTACAACGA	TCCCAAGC		GCTTGGGA A	ACCACGGG	
1254	CTGCGCGT	GGCTAGCTACAACGA	GGTTCCCA		TGGGAACC A	ACGCGCAG	
1256	CACTGCGC	GGCTAGCTACAACGA	GTGGTTCC		GGAAACCAC G	GCGCAGTG	
1258	GGCACTGC	GGCTAGCTACAACGA	GCGTGGTT		AACCACGC G	GCAGTGCC	
1261	AGGGGCAC	GGCTAGCTACAACGA	TGCGCGTG		CACGCGCA G	GTGCCCTT	
1263	GTAGGGGC	GGCTAGCTACAACGA	ACTGCGCG		CGCGCAGT G	GCCCCCTAC	
1269	CACCCCGT	GGCTAGCTACAACGA	AGGGGCAC		GTGCCCTT A	ACGGGGTG	
1274	AGGAGCAC	GGCTAGCTACAACGA	CCCGTAGG		CCTACGGG G	GTGCTCCT	
1276	TGAGGAGC	GGCTAGCTACAACGA	ACCCCGTA		TACGGGGT G	GCTCCTCA	
1286	CAGTGCGT	GGCTAGCTACAACGA	CTTGAGGA		TCCTCAAG A	ACGCACTG	
1288	GGCAGTGC	GGCTAGCTACAACGA	GTCTTGAG		CTCAAGAC G	GCACTGCC	
1290	CGGGCAGT	GGCTAGCTACAACGA	GCGTCTTG		CAAGACGC A	ACTGCCCG	
1293	CAGCGGGC	GGCTAGCTACAACGA	AGTGCGTC		GACGCACT G	GCCCCGCTG	
1297	CTCGCAGC	GGCTAGCTACAACGA	GGGCAGTG		CAC TGCCC G	GCTGCGAG	
1300	CAGCTCGC	GGCTAGCTACAACGA	AGCGGGCA		TGCCCGCT G	GCGAGCTG	
1304	ACCGCAGC	GGCTAGCTACAACGA	TGCGAGCG		CGCTGCGA G	GCTGCGGT	
1307	GTGACCGC	GGCTAGCTACAACGA	AGCTCGCA		TGCGAGCT G	GCGGTCCAC	
1310	GGGGTGAC	GGCTAGCTACAACGA	GCGAGTTC		GAGTGGG G	GTACACCC	
1313	GCTGGGGT	GGCTAGCTACAACGA	GACCGCAG		CTGCGGTC A	ACCCACGC	
1319	CCGGCTGC	GGCTAGCTACAACGA	TGGGGTGA		TCACCCCA G	GCAGCCGG	
1322	ACACCCGC	GGCTAGCTACAACGA	TGCTGGGG		CCCCAGCA G	GCCGGTGT	
1326	ACAGACAC	GGCTAGCTACAACGA	CGGCTGCT		AGCAGCCG G	GTGTCTGT	
1328	GCACAGAC	GGCTAGCTACAACGA	ACCGGCTG		CAGCCGGT G	GTCTGTGC	
1332	CCGGGCAC	GGCTAGCTACAACGA	AGACACCG		CGGTGTCT G	GTGCCCGG	
1334	TCCCGGGC	GGCTAGCTACAACGA	ACAGACAC		GTGTCTGT G	GCCCCGGA	
1345	CCTGGGGC	GGCTAGCTACAACGA	TTCTCCCG		CGGGAGAA G	GCCCCAGG	
1353	CACAGAGC	GGCTAGCTACAACGA	CCTGGGCG		GCCCCAGG G	GCTCTGTG	
1358	GCCGCCAC	GGCTAGCTACAACGA	AGAGCCCT		AGGCTCT G	GTGGGGCC	
1361	GGGGCCGC	GGCTAGCTACAACGA	CACAGAGC		GCTCTGTG G	GCGGCCCC	
1364	TGCGGGGC	GGCTAGCTACAACGA	GGCCACAG		CTGTGGCG G	GCCCCGGA	

Table 16

1380	GTCTGTGT	GGCTAGCTACAACGA	CCTCTCTCC		GGAGGAGG A ACACAGAC	
1382	GGTCTGT	GGCTAGCTACAACGA	GTCTCTCT		AGGAGGAC A ACAGACCC	
1386	ACGGGGGT	GGCTAGCTACAACGA	CTGTGTCC		GGACACAG A ACCCCCGT	
1392	CAGGCGAC	GGCTAGCTACAACGA	GGGGGTCT		AGACCCCC G GTCGCCCTG	
1395	CACCAGGC	GGCTAGCTACAACGA	GACGGGGG		CCCCCGTC G GCCTGGTG	
1400	AGCTGCAC	GGCTAGCTACAACGA	CAGGCGAC		GTGCGCTG G GTGCAGCT	
1402	GCAGCTGC	GGCTAGCTACAACGA	ACCAGGCG		CGCCTGGT G GCAGCTGC	
1405	GGAGCAGC	GGCTAGCTACAACGA	TGCACCCAG		CTGGTGCA G GCTGCTCC	
1408	GGCGGAGC	GGCTAGCTACAACGA	AGCTGCAC		GTGCAGCT G GCTCCGCC	
1413	GTGCTGGC	GGCTAGCTACAACGA	GGAGCAGC		GCTGCTCC G GCCAGCAC	
1417	TGCTGTGC	GGCTAGCTACAACGA	TGGCGGAG		CTCCGCCA G GCACAGCA	
1419	GCTGCTGT	GGCTAGCTACAACGA	GCTGGCGG		CCGCCAGC A ACAGCAGC	
1422	GGGGCTGC	GGCTAGCTACAACGA	TGTGCTGG		CCAGCACA G GCAGCCCC	
1425	CCAGGGGC	GGCTAGCTACAACGA	TGCTGTGC		GCACAGCA G GCCCCTGG	
1432	ACACCTGC	GGCTAGCTACAACGA	CAGGGGCT		AGCCCTTG G GCAGGTGT	
1436	CCGTACAC	GGCTAGCTACAACGA	CTGCCAGG		CCTGGCAG G GTGTACGG	
1438	AGCCGTAC	GGCTAGCTACAACGA	ACCTGCCA		TGGCAGGT G GTACGGCT	
1440	GAAGCCGT	GGCTAGCTACAACGA	ACACCTGC		GCAGGTGT A ACGGCTTC	
1443	CACGAAGC	GGCTAGCTACAACGA	CGTACACC		GGTGTACG G GCTTCGTG	
1448	GCCCGCAC	GGCTAGCTACAACGA	GAAGCCGT		ACGGCTTC G GTGCGGGC	
1450	AGGCCCGC	GGCTAGCTACAACGA	ACGAAGCC		GGCTTCGT G GCGGGCCT	
1454	AGGCAGGC	GGCTAGCTACAACGA	CCGCACGA		TCGTGCGG G GCCTGCCT	
1458	GCGCAGGC	GGCTAGCTACAACGA	AGGCCCGC		GCGGGCCT G GCCTGCGC	
1462	GCCGGCGC	GGCTAGCTACAACGA	AGGCAGGC		GCCTGCCT G GCGCCGCG	
1464	CAGCCGGC	GGCTAGCTACAACGA	GCAGGCGAG		CTGCTGCG G GCCGGCTG	
1468	GCACCCAGC	GGCTAGCTACAACGA	CGGCGCAG		CTGCGCCG G GCTGGTGC	
1472	GGGGGCAC	GGCTAGCTACAACGA	CAGCCGCG		GCCGGCTG G GTGCCCCC	
1474	CTGGGGGC	GGCTAGCTACAACGA	ACCAGCCG		CGGCTGGT G GCCCCACG	
1482	CCAGAGGC	GGCTAGCTACAACGA	CTGGGGGC		GCCCCCAG G GCCTCTGG	
1491	CCTGGAGC	GGCTAGCTACAACGA	CCCAGAGG		CCTCTGGG G GCTCCAGG	
1498	CGTTGTGC	GGCTAGCTACAACGA	CTGGAGCC		GGCTCCAG G GCACAACG	

Table 16

1500	TTGTTGT	GGCTAGCTACAACGA	GCTGGAG		CTCCAGGC A ACAACGAA
1503	GGTTTCGT	GGCTAGCTACAACGA	TGTGCCTG		CAGGCACA A ACGAACGC
1507	AGCGGCGT	GGCTAGCTACAACGA	TGTTGTG		CACAACGA A ACGCCGCT
1509	GAAGCGGC	GGCTAGCTACAACGA	GTTCTGTG		CAACGAAC G GCCGCTTC
1512	GAGGAAGC	GGCTAGCTACAACGA	GGGTTTCG		CGAACGCC G GCTTCCTC
1524	CTTGGTGT	GGCTAGCTACAACGA	TCCTGAGG		CCTCAGGA A ACACCAAG
1526	TTCTTGGT	GGCTAGCTACAACGA	GTTCTCTGA		TCAGGAAC A ACCAAGAA
1534	AGATGAAC	GGCTAGCTACAACGA	TTCTTGGT		ACCAAGAA G GTTCATCT
1538	AGGGAGAT	GGCTAGCTACAACGA	GAATTTCT		AGAATTTC A ATCTCCCT
1552	TGGCATGC	GGCTAGCTACAACGA	TTCCCCAG		CTGGGAA G GCATGCCA
1554	CTTGGCAT	GGCTAGCTACAACGA	GCTTCCCC		GGGAAGC A ATGCCAAG
1556	AGCTTGGC	GGCTAGCTACAACGA	ATGCTTCC		GGAGCAT G GCCAAGCT
1561	GGGAGAGC	GGCTAGCTACAACGA	TTGGCATG		CATGCCAA G GCTCTCGC
1567	CCTGCAGC	GGCTAGCTACAACGA	GAGAGCTT		AAGCTCTC G GCTGCAGG
1570	GCTCCTGC	GGCTAGCTACAACGA	AGCGAGAG		CTCTCGCT G GCAGGAGC
1576	AGTCAGC	GGCTAGCTACAACGA	TCCTGCAG		CTGCAGGA G GCTGACGT
1580	TTCCACGT	GGCTAGCTACAACGA	CAGTCTCT		AGGAGCTG A ACGTGGAA
1582	TCTTCCAC	GGCTAGCTACAACGA	GTCAGCTC		GAGCTGAC G GTGGAAGA
1589	ACGCTCAT	GGCTAGCTACAACGA	CTTCCACG		CGTGAAG A ATAGCGT
1593	CCGCACGC	GGCTAGCTACAACGA	TCATCTTC		GAAGATGA G GCGTGC GG
1595	TCCCGCAC	GGCTAGCTACAACGA	GCTCATCT		AGATGAGC G GTGCGGGA
1597	AGTCCCGC	GGCTAGCTACAACGA	ACGCTCAT		ATGAGCGT G GCGGAGCT
1602	AGCGCAGT	GGCTAGCTACAACGA	CCCGCAGG		CGTGC GGG A ACTGCGCT
1605	CCAAGCGC	GGCTAGCTACAACGA	AGTCCCGC		GCGGAGCT G GCGCTTGG
1607	AGCCAAGC	GGCTAGCTACAACGA	GCAGTCCC		GGGACTGC G GCTTGGCT
1612	TGCGCAGC	GGCTAGCTACAACGA	CAAGCGCA		TGCGCTTG G GCTGCGCA
1615	TCCTGCGC	GGCTAGCTACAACGA	AGCCAAGC		GCTTGGCT G GCGCAGGA
1617	GCTCTGTC	GGCTAGCTACAACGA	GCAGCCAA		TTGGCTGC G GCAGGAGC
1623	CCCTGGGC	GGCTAGCTACAACGA	TCCTGCGC		GGCAGGA G GCCAGGG
1631	CAGCCAAC	GGCTAGCTACAACGA	CCCTGGGC		GCCAGGG G GTTGGCTG
1635	AACACAGC	GGCTAGCTACAACGA	CAACCCCT		AGGGTTG G GCTGTGTT

Table 16

1638	CGGAACAC	GGCTAGCTACAACGA	AGCCAACC		GGTTGGCT G GTGTTCCG
1640	GCCGGAAC	GGCTAGCTACAACGA	ACAGCCAA		TTGGCTGT G GTTCCGGC
1646	TCTGCGGC	GGCTAGCTACAACGA	CGGAACAC		GTGTTCCG G GCGCAGA
1649	TGCTCTGC	GGCTAGCTACAACGA	GGCGGAA		TTCCGGCC G GCAGAGCA
1654	GACGGTGC	GGCTAGCTACAACGA	TCTGCGGC		GCGCAGA G GCACCCTC
1656	CAGACGGT	GGCTAGCTACAACGA	GCTCTGGC		CGCAGAGC A ACCGTCTG
1659	ACGCAGAC	GGCTAGCTACAACGA	GGTGTCTT		AGAGCAC C G GTCTGCGT
1663	CCTCACGC	GGCTAGCTACAACGA	AGACGGTG		CACCGTCT G GCGTGAGG
1665	CTCCTCAC	GGCTAGCTACAACGA	GCAGACGG		CCGTCTGC G GTGAGGAG
1673	GCCAGGAT	GGCTAGCTACAACGA	CTCCTCAC		GTGAGGAG A ATCCTGGC
1679	AACTTGGC	GGCTAGCTACAACGA	CAGGATCT		AGATCCTG G GCCAAGTT
1684	GCAGGAAC	GGCTAGCTACAACGA	TTGGCCAG		CTGGCCAA G GTTCTGTC
1690	GCCAGTGC	GGCTAGCTACAACGA	AGGAATTT		AAGTTCTT G GCACTGGC
1692	CAGCCAGT	GGCTAGCTACAACGA	GCAGGAAC		GTTCCTGC A ACTGGCTG
1696	TCATCAGC	GGCTAGCTACAACGA	CAGTGCAG		CTGCACTG G GCTGATGA
1700	ACACTCAT	GGCTAGCTACAACGA	CAGCCAGT		ACTGGCTG A ATGAGTGT
1704	GTACACAC	GGCTAGCTACAACGA	TCATCAGC		GCTGATGA G GTGTGTAC
1706	ACGTACAC	GGCTAGCTACAACGA	ACTCATCA		TGATGAGT G GTGTACGT
1708	CGACGTAC	GGCTAGCTACAACGA	ACACTCAT		ATGAGTGT G GTACGTCC
1710	GAGGACGT	GGCTAGCTACAACGA	ACACACTC		GAGTGTGT A ACGTCGTC
1712	TCGACGAC	GGCTAGCTACAACGA	GTACACAC		GTGTGTAC G GTCGTGCA
1715	AGCTCGAC	GGCTAGCTACAACGA	GACGTACA		TGTACGTC G GTCGAGCT
1720	TGAGCAGC	GGCTAGCTACAACGA	TGACGAC		GTCGTGCA G GCTGTCTCA
1723	ACCTGAGC	GGCTAGCTACAACGA	AGCTCGAC		GTCGAGCT G GCTCAGGT
1729	AGAAAGAC	GGCTAGCTACAACGA	CTGAGCAG		CTGCTCAG G GTCCTTCT
1740	CGTGACAT	GGCTAGCTACAACGA	AAAAGAAA		TTTCTTTT A ATGTCACG
1742	TCGGTGAC	GGCTAGCTACAACGA	ATAAAAGA		TCTTTTAT G GTCACGGA
1745	GTCTCCGT	GGCTAGCTACAACGA	GACATAAA		TTTATGTC A ACGGAGAC
1751	AACGTGGT	GGCTAGCTACAACGA	CTCCGTGA		TCACGGAG A ACCACGTT
1754	TGAACAGT	GGCTAGCTACAACGA	GGTCTCCG		CGGAGACC A ACGTTTCA
1756	TTTGA AAC	GGCTAGCTACAACGA	GTGGTCTC		GAGACCAC G GTTTCAAA

Table 16

1767	GAGCTGT GGCTAGCTACAACGA TCTTTGA		TCAAAGA A ACAGGCTC
1771	AAAAGAGC GGCTAGCTACAACGA CTGTTCTT		AAGAACAG G GCTCTTTT
1782	CTTCCGGT GGCTAGCTACAACGA AGAAAAAG		CTTTTCT A ACCGGAAG
1791	CCAGACAC GGCTAGCTACAACGA TCTTCCGG		CCGGAAGA G GTGTCTGG
1793	CTCCAGAC GGCTAGCTACAACGA ACTCTTCC		GGAAGAGT G GTCTGGAG
1800	CAACTTGC GGCTAGCTACAACGA TCCAGACA		TGCTTGA G GCAAGTTG
1804	TTTGCAAC GGCTAGCTACAACGA TTGCTCCA		TGGAGCA G GTTGCAAA
1807	TGCTTTGC GGCTAGCTACAACGA AACTTGCT		AGCAAGTT G GCAAAGCA
1812	TCCAATGC GGCTAGCTACAACGA TTTGCAAC		GTTCAAA G GCATTGGA
1814	ATTCCAAT GGCTAGCTACAACGA GCTTTGCA		TGCAAGC A ATTGGAAT
1820	TGCTGTAT GGCTAGCTACAACGA TCCAATGC		GCATTGA A ATCAGACA
1825	AGTGCTGT GGCTAGCTACAACGA CTGATTCC		GGAAATCAG A ACAGCACT
1828	TCAAGTGC GGCTAGCTACAACGA TGTCTGAT		ATCAGACA G GCACTTGA
1830	CTTCAAGT GGCTAGCTACAACGA GCTGTCTG		CAGACAGC A ACTTGAAG
1841	AGCTGCAC GGCTAGCTACAACGA CTTCTTCA		TGAAGAG G GTGCAGCT
1843	GCAGCTGC GGCTAGCTACAACGA ACCCTCTT		AAGAGGT G GCAGCTGC
1846	CCCGCAGC GGCTAGCTACAACGA TGCACCCCT		AGGTGCA G GCTGCGGG
1849	GCTCCCGC GGCTAGCTACAACGA AGCTGCAC		GTGCAGCT G GCGGAGC
1855	CCGACAGC GGCTAGCTACAACGA TCCCGCAG		CTGCGGA G GCTGTGG
1858	CTTCCGAC GGCTAGCTACAACGA AGCTCCCG		CGGAGCT G GTCGGAAG
1865	ACCTCTGC GGCTAGCTACAACGA TTCCGACA		TGTCGGA G GCAGAGGT
1871	TGCCCTGAC GGCTAGCTACAACGA CTCTGCTT		AAGCAGAG G GTCAGGCA
1876	GATGCTGC GGCTAGCTACAACGA CTGACCTC		GAGGTCA G GCAGCATC
1879	CCCGATGC GGCTAGCTACAACGA TGCTGAC		GTCAAGCA G GCATCGGG
1881	TTCCCCGAT GGCTAGCTACAACGA GCTGCTG		CAGGAGC A ATCGGGA
1889	GGCTGGC GGCTAGCTACAACGA TTCCCGAT		ATCGGAA G GCCAGGCC
1894	GGGCGGC GGCTAGCTACAACGA CTGGCTTC		GAAGCCAG G GCCCGCCC
1898	AGCAGGGC GGCTAGCTACAACGA GGGCTGG		CCAGGCC G GCCCTGCT
1903	ACGTCAGC GGCTAGCTACAACGA AGGGCGGG		CCCGCCT G GCTGACGT
1907	CTGGACGT GGCTAGCTACAACGA CAGCAGGG		CCCTGCTG A AGTCCAG
1909	GTCTGGAC GGCTAGCTACAACGA GTGAGCAG		CTGCTGAC G GTCAGAGC

Table 16

1915	AGCGGAGT	GGCTAGCTACAACGA	CTGGACGT		ACGTCAG A	ACTCCGCT
1920	GATGAAGC	GGCTAGCTACAACGA	GGAGTCTG		CAGACTCC G	GCTTCATC
1925	TTGGGGAT	GGCTAGCTACAACGA	GAACGGGA		TCCGCTTC A	ATCCCCAA
1933	CGTCAGGC	GGCTAGCTACAACGA	TTGGGGAT		ATCCCCAA G	GCCTGACG
1938	CAGCCCGT	GGCTAGCTACAACGA	CAGGCTTG		CAAGCCTG A	ACGGGCTG
1942	GCCGCAGC	GGCTAGCTACAACGA	CCGTCAAG		CCTGACGG G	GCTGCGGC
1945	TCGGCCGC	GGCTAGCTACAACGA	AGCCGGTC		GACGGGCT G	GGGGCCGA
1948	CAATCGGC	GGCTAGCTACAACGA	CGCAGCCC		GGGCTGCG G	GCGGATTG
1952	TTCAACAAT	GGCTAGCTACAACGA	CGGGCGCA		TGCGGCCG A	ATTGTGAA
1955	ATGTTTAC	GGCTAGCTACAACGA	AATCGGCC		GGCCGATT G	GTGAACAT
1959	GTCCATGT	GGCTAGCTACAACGA	TCACAATC		GATTGTGA A	ACATGGAC
1961	TAGTCCAT	GGCTAGCTACAACGA	GTTCACAA		TTGTGAAC A	ATGGACTA
1965	GACGTAGT	GGCTAGCTACAACGA	CCATGTTT		GAACATGG A	ACTACGTC
1968	CACGACGT	GGCTAGCTACAACGA	AGTCCATG		CATGGACT A	ACGTCGTG
1970	CCCACGAC	GGCTAGCTACAACGA	GTAATCCA		TGGACTAC G	GTGTTGGG
1973	GCTCCAC	GGCTAGCTACAACGA	GACGTAGT		ACTACGTC G	GTGGGAGC
1979	GTTCTGGC	GGCTAGCTACAACGA	TCCCACGA		TCGTGGGA G	GCCAGAAC
1985	CGGAACGT	GGCTAGCTACAACGA	TCTGGCTC		GAGCCAGA A	ACGTTCCG
1987	TGCGGAAC	GGCTAGCTACAACGA	GTTCTGGC		GCCAGAAC G	GTTCGCGA
1992	TTCTCTGC	GGCTAGCTACAACGA	GGAGGTT		AACGTTCC G	GCAGAGAA
2006	CGTCGGC	GGCTAGCTACAACGA	CCTCTTTT		AAAAGAGG G	GCCGAGCG
2011	TGAGACGC	GGCTAGCTACAACGA	TCGGCCCT		AGGGCCGA G	GCGTCTCA
2013	GGTGAGAC	GGCTAGCTACAACGA	GCTCGGCC		GGCCGAGC G	GTCTCACC
2018	CTCGAGGT	GGCTAGCTACAACGA	GAGACGCT		AGCGTCTC A	ACCTGAG
2027	GCCTTCAC	GGCTAGCTACAACGA	CCTCGAGG		CCTCGAGG G	GTGAAGGC
2033	AACAGTGC	GGCTAGCTACAACGA	CTTCACCC		GGGTGAAG G	GCACTGTT
2035	TGAACAGT	GGCTAGCTACAACGA	GCCTTCAC		GTGAAGGC A	ACTGTTCA
2038	CGCTGNAC	GGCTAGCTACAACGA	AGTGCCTT		AAGGCACT G	GTTCAGCG
2043	GAGCACGC	GGCTAGCTACAACGA	TGAACAGT		ACTGTTCA G	GCGTCTC
2045	TTGAGCAC	GGCTAGCTACAACGA	GCTGAACA		TGTTTCAGC G	GTGCTCAA
2047	AGTTGAGC	GGCTAGCTACAACGA	ACGCTGAA		TTTACGCGT G	GCTCAACT

Table 16

2052	CTCGTAGT	GGCTAGCTACAACGA	TGAGCAG	CGTGCTCA A	ACTACGAG
2055	CCGTCGT	GGCTAGCTACAACGA	AGTTGAGC	GCTCACT A	ACGAGCGG
2059	GGCCCGC	GGCTAGCTACAACGA	TCGTAGTT	AATACGA G	GCGGGCGC
2063	CGCCGCG	GGCTAGCTACAACGA	CGCTCGT	ACGAGCG G	GCGGGCGG
2065	GGCGCGC	GGCTAGCTACAACGA	GCCGCTC	GAGCGGC G	GCGGCGCC
2068	CGGGCGC	GGCTAGCTACAACGA	CGCGCCG	CGGCGCG G	GCGCCCGG
2070	GCCGGGC	GGCTAGCTACAACGA	GCCGGCC	GCGCGCG G	GCGCCCGG
2076	CAGGAGC	GGCTAGCTACAACGA	CGGGGCG	GCGCCCG G	GCCTCCTG
2085	AGAGGCG	GGCTAGCTACAACGA	CCAGGAG	CCTCCTG G	GCGCCTCT
2087	ACAGGAC	GGCTAGCTACAACGA	GCCAGGA	TCCTGGC G	GCCTCTGT
2093	CCCGGAG	GGCTAGCTACAACGA	ACAGAGC	GCGCCTCT G	GTGCTGGG
2095	GGCCAGC	GGCTAGCTACAACGA	ACAGAGC	GCCTCTGT G	GCTGGGCC
2100	GTCCAGC	GGCTAGCTACAACGA	CCAGCACA	TGTGCTGG G	GCCTGGAC
2106	GATATCGT	GGCTAGCTACAACGA	CCAGGCC	GGCCTGG A	ACGATATC
2109	GTGGATAT	GGCTAGCTACAACGA	GTCCAGG	CCTGGAGC A	ATATCCAC
2111	CTGTGGAT	GGCTAGCTACAACGA	ATGTCGA	TGGACGAT A	ATCCACAG
2115	GGCCCTGT	GGCTAGCTACAACGA	GGATATCG	CGATATCC A	ACAGGGCC
2120	CGCCAGC	GGCTAGCTACAACGA	CCTGTGGA	TCCACAGG G	GCCTGGCG
2125	AGGTGCG	GGCTAGCTACAACGA	CAGGCCCT	AGGCGCTG G	GCGCACCT
2127	GAAGGTG	GGCTAGCTACAACGA	GCCAGGCC	GCGCTGGC G	GCACCTTC
2129	ACGAAGT	GGCTAGCTACAACGA	GCGCCAGG	CCTGGCGC A	ACCTTCGT
2135	CGCAGAC	GGCTAGCTACAACGA	GAAGGTGC	GCACCTTC G	GTGCTGGG
2137	CACGCAGC	GGCTAGCTACAACGA	ACGAAGGT	ACCTTCGT G	GCTGCGTG
2140	GCACACGC	GGCTAGCTACAACGA	AGCACGAA	TTCGTGCT G	GCGTGTGC
2142	CCGCACAC	GGCTAGCTACAACGA	GCAGCACG	CGTGCTGC G	GTGTGCGG
2144	GCCCGCAC	GGCTAGCTACAACGA	ACGCAGCA	TGCTGCGT G	GTGCGGGC
2146	GGCCCGGC	GGCTAGCTACAACGA	ACACGCAG	CTGCTGT G	GCGGGGCC
2150	TCCTGGGC	GGCTAGCTACAACGA	CCGCACAC	GTGTGGG G	GCCCAGGA
2157	CGCGGGT	GGCTAGCTACAACGA	CCTGGGCC	GGCCAGG A	ACCGCGG
2161	CAGCGGC	GGCTAGCTACAACGA	GCGTCCTG	CAGGACC G	GCCGCTG
2164	GCTCAGC	GGCTAGCTACAACGA	GCGGGTGC	GACCCGC G	GCCTGAGC

Table 16

2170	AGTACAGC	GGCTAGCTACAACGA	TCAGGCGG		CCGCCCTGA	G	GCTGTACT
2173	CAAAGTAC	GGCTAGCTACAACGA	AGCTCAGG		CCTGAGCT	G	GTACTTTG
2175	GACAAAGT	GGCTAGCTACAACGA	ACAGCTCA		TGAGCTGT	A	ACTTTGTC
2180	ACCTTGAC	GGCTAGCTACAACGA	AAAGTACA		TGTACTTT	G	GTCAAGGT
2186	ACATCCAC	GGCTAGCTACAACGA	CTTGACAA		TTGTCAAG	G	GTGGATGT
2190	CGTCACAT	GGCTAGCTACAACGA	CCACCTTG		CAAGGTGG	A	ATGTGACG
2192	CCCGTCAC	GGCTAGCTACAACGA	ATCCACCT		AGGTGGAT	G	GTACGGG
2195	GCGCCCGT	GGCTAGCTACAACGA	CACATCCA		TGGATGTG	A	ACGGGCGC
2199	GTACGGC	GGCTAGCTACAACGA	CGGTACCA		TGTGACGG	G	GGCGGTAC
2201	TCGTACGC	GGCTAGCTACAACGA	GCCGTCA		TGACGGGC	G	GGTACGGA
2203	TGTCGTAC	GGCTAGCTACAACGA	GCGCCGT		ACGGGCGC	G	GTACGACA
2205	GGTGTGCT	GGCTAGCTACAACGA	ACGGCCCC		GGGCGCGT	A	ACGACACC
2208	GATGCTGT	GGCTAGCTACAACGA	CGTAGCGG		CGGTACG	A	ACACCATC
2210	GGGATGGT	GGCTAGCTACAACGA	GTCGTACG		CGTAGGAC	A	ACCATCCC
2213	TGGGGGAT	GGCTAGCTACAACGA	GGTGTGCT		ACGACACC	A	ATCCCCCA
2223	GAGCCTGT	GGCTAGCTACAACGA	CCTGGGGG		CCCCCAGG	A	ACAGGCTC
2227	CCGTGAGC	GGCTAGCTACAACGA	CTGTCTGT		CAGGACAG	G	GCTCACGG
2231	ACCTCCGT	GGCTAGCTACAACGA	GAGCCTGT		ACAGGCTC	A	ACGGAGGT
2237	GCGATGAC	GGCTAGCTACAACGA	CTCGTGA		TCACGGAG	G	GTCAATGC
2240	CTGGCGAT	GGCTAGCTACAACGA	GACCTCCG		CGGAGGTC	A	ATCGCCAG
2243	ATGCTGGC	GGCTAGCTACAACGA	GATGACCT		AGGTCATC	G	GCCAGCAT
2247	GATGATGC	GGCTAGCTACAACGA	TGGCGATG		CATCGCCA	G	GCATCATC
2249	TTGATGAT	GGCTAGCTACAACGA	GCTGGCGA		TGCGCAGC	A	ATCATCAA
2252	GGTTTGAT	GGCTAGCTACAACGA	GATGCTGG		CCAGCATC	A	ATCAAACC
2257	TCTGGGGT	GGCTAGCTACAACGA	TTGATGAT		ATCATCAA	A	ACCCAGA
2265	GTACGTGT	GGCTAGCTACAACGA	TCTGGGGT		ACCCAGA	A	ACAGGTAC
2267	CAGTAGGT	GGCTAGCTACAACGA	GTTCTGGG		CCCAGAAC	A	ACGTACTG
2269	CGCAGTAC	GGCTAGCTACAACGA	GTGTTCTG		CAGAACAC	G	GTACTGCG
2271	CACGCAGT	GGCTAGCTACAACGA	ACGTGTTT		GNACACGT	A	ACTGGGTG
2274	ACGCACGC	GGCTAGCTACAACGA	AGTAGCTG		CAGGTACT	G	GGGTGCGT
2276	CGACGCAC	GGCTAGCTACAACGA	GCAGTACG		CGTACTGC	G	GTGGGTG

Table 16

2278	ACGAGCG	GGCTAGCTACAACGA	ACGCAGTA		TACTGCGT G GCGTCGGT
2280	ATACCGAC	GGCTAGCTACAACGA	GCACGCAG		CTGCGTGC G GTCGGTAT
2284	CGGCATAC	GGCTAGCTACAACGA	CGACGCAC		GTGCGTCG G GTATGCCG
2286	CACGGCAT	GGCTAGCTACAACGA	ACCGAGCG		GCGTCGGT A ATGCCGTG
2288	ACCACGGC	GGCTAGCTACAACGA	ATACCGAC		GTCGGTAT G GCCGTGGT
2291	TGGACCCAC	GGCTAGCTACAACGA	GGCATATAC		GGTATGCC G GTGTCGA
2294	TTCTGGAC	GGCTAGCTACAACGA	CACGGCAT		ATGCCGTG G GTCAGAA
2303	TGGCGGC	GGCTAGCTACAACGA	CTTCTGGA		TCCAGAAG G GCCGCCA
2306	CCATGGGC	GGCTAGCTACAACGA	GGCTTTCT		AGAAGGCC G GCCATGG
2310	GTGCCCAT	GGCTAGCTACAACGA	GGCGGCC		GGCGGCC A ATGGCAC
2314	GGACGTGC	GGCTAGCTACAACGA	CCATGGGC		GCCCATGG G GCACGTCC
2316	GCGGACGT	GGCTAGCTACAACGA	GCCCATGG		CCATGGGC A ACGTCCGC
2318	TTGCGGAC	GGCTAGCTACAACGA	GTGCCCAT		ATGGGCAC G GTCGCAA
2322	GGCCTTGC	GGCTAGCTACAACGA	GGACGTGC		GCACGTCC G GCAAGGCC
2327	TTGAAGGC	GGCTAGCTACAACGA	CTTGCGGA		TCCGAAG G GCCTTCAA
2337	GACGTGGC	GGCTAGCTACAACGA	TCTTGAAG		CTTCAAGA G GCCACGTC
2340	AGAGACGT	GGCTAGCTACAACGA	GGCTCTTG		CAAGAGCC A ACGTCTCT
2342	GTAGAGAC	GGCTAGCTACAACGA	GTGGCTCT		AGAGCCAC G GTCTCTAC
2348	GTCAAGGT	GGCTAGCTACAACGA	AGAGACGT		ACGTCTCT A ACCTTGAC
2354	AGGTCTGT	GGCTAGCTACAACGA	CAAGGTAG		CTACCTTG A ACAGACCT
2358	CTGGAGGT	GGCTAGCTACAACGA	CTGTCAAG		CTTGACAG A ACCTCCAG
2365	TGTACGGC	GGCTAGCTACAACGA	TGGAGGTC		GACCTCCA G GCCGTACA
2368	GCATGTAC	GGCTAGCTACAACGA	GGCTGGAG		CTCCAGCC G GTACATGC
2370	TGCGCATGT	GGCTAGCTACAACGA	ACGGCTGG		CCAGCGT A ACATGCGA
2372	TGTCGCAT	GGCTAGCTACAACGA	GTACGGCT		AGCCGTAC A ATGGACA
2374	ACTGTGCG	GGCTAGCTACAACGA	ATGTACGG		CCGTACAT G GCGACAGT
2377	CGAACTGT	GGCTAGCTACAACGA	CGCATGTA		TACATGCG A ACAGTTCC
2380	CCACGAAC	GGCTAGCTACAACGA	TGTGCGAT		ATGCGACA G GTTCGTGG
2384	TGAGCCAC	GGCTAGCTACAACGA	GAACTGTC		GACAGTTC G GTGGCTCA
2387	AGGTGAGC	GGCTAGCTACAACGA	CACGAAC		AGTTCGTG G GCTCACCT
2391	CTGCAGGT	GGCTAGCTACAACGA	GAGCCACG		GCTGGCTC A ACCTGCAG

Table 16

2395	TCTCTGCG	GGCTAGCTACAACGA	AGGTGAGC		GCTCACCT G GCAGGAGA
2402	GGGCTGGT	GGCTAGCTACAACGA	CTCCTGCA		TGCAGGAG A ACCAGCCC
2406	CAGCGGGC	GGCTAGCTACAACGA	TGGTCTCC		GGAGACCA G GCCCGCTG
2410	CCCTCAGC	GGCTAGCTACAACGA	GGGCTGGT		ACCAGCCC G GCTGAGGG
2418	GACGGCAT	GGCTAGCTACAACGA	CCCTCAGC		GCTGAGGG A ATGCCGTC
2420	ACGACGGC	GGCTAGCTACAACGA	ATCCCTCA		TGAGGGAT G GCCGTCGT
2423	ATGACGAC	GGCTAGCTACAACGA	GGCATCCC		GGGATGCC G GTCGTCTAT
2426	TCGATGAC	GGCTAGCTACAACGA	GACGGCAT		ATGCCGTC G GTCATCGA
2429	TGCTCGAT	GGCTAGCTACAACGA	GAGCAAGG		CCGTCTGC A ATCGAGCA
2434	AGCTCTGC	GGCTAGCTACAACGA	TCGATGAC		GTCATCGA G GCAGAGCT
2439	GGAGGAGC	GGCTAGCTACAACGA	TCTGCTCG		CGAGCAGA G GCTCCTCC
2451	GGCCTCAT	GGCTAGCTACAACGA	TCAGGGAG		CTCCCTGA A ATGAGGCC
2456	CTGCTGGC	GGCTAGCTACAACGA	CTCATTTA		TGAATGAG G GCCAGCAG
2460	GCCACTGC	GGCTAGCTACAACGA	TGGCCTCA		TGAGGCCA G GCAGTGGC
2463	GAGGCCAC	GGCTAGCTACAACGA	TGCTGGCC		GGCCAGCA G GTGGCCTC
2466	GAAGAGGC	GGCTAGCTACAACGA	CACCTGCTG		CAGCAGTG G GCCTCTTC
2475	GAAGACGT	GGCTAGCTACAACGA	CGAAGAGG		CCTCTTGG A ACGTCTTC
2477	AGGAAGAC	GGCTAGCTACAACGA	GTGGAAGA		TCTTCGAC G GTCTTCCT
2485	TGAAGCGT	GGCTAGCTACAACGA	AGGAAGAC		GTCTTCCT A ACGCTTCA
2487	CATGAAGC	GGCTAGCTACAACGA	GTAGGAAG		CTTCCTAC G GCTTCATG
2492	TGGCACAT	GGCTAGCTACAACGA	GAAGCGTA		TAGGCTTC A ATGTGCCA
2494	GGTGGCAC	GGCTAGCTACAACGA	ATGAAGCG		CGCTTCAT G GTGCCACC
2496	GTGTGGC	GGCTAGCTACAACGA	ACATGAAG		CTTCATGT G GCCACCAC
2499	GGCGTGGT	GGCTAGCTACAACGA	GGCACATG		CATGTGCC A ACCAGGCC
2502	CACGGCGT	GGCTAGCTACAACGA	GGTGGCAC		GTGGCCAC A ACGCGGTG
2504	CGCACGGC	GGCTAGCTACAACGA	GTGTGGC		GCCACCAC G GCCGTGCG
2507	ATGCGCAC	GGCTAGCTACAACGA	GGCGTGGT		ACCACGCC G GTGCGCAT
2509	TGATGCGC	GGCTAGCTACAACGA	ACGGCGTG		CACGCCGT G GCGCATCA
2511	CCTGATGC	GGCTAGCTACAACGA	GCACGGCG		CGCCGTGC G GCATCAGG
2513	CCCCTGAT	GGCTAGCTACAACGA	GCGCAGCG		CGGTGGC A ATCAGGGG
2520	GGACTTGC	GGCTAGCTACAACGA	CCCTGATG		CATCAGGG G GCAGTCC

Table 16

2524	CGTAGGAC	GGCTAGCTACAACGA	TTGCCCT		AGGGGCAA	G	GTCTACG
2529	CTGGACGT	GGCTAGCTACAACGA	AGGACTTG		CAAGTCCT	A	ACGTCCAG
2531	CACTGGAC	GGCTAGCTACAACGA	GTAGACT		AGTCCTAC	G	GTCCAGTG
2536	CCTGGCAC	GGCTAGCTACAACGA	TGGACGTA		TACGTCCA	G	GTGCCAGG
2538	CCCCTGGC	GGCTAGCTACAACGA	ACTGGACG		CGTCCAGT	G	GCCAGGGG
2546	TGCGGGAT	GGCTAGCTACAACGA	CCCCTGGC		GCCAGGGG	A	ATCCGGCA
2551	AGCCCTGC	GGCTAGCTACAACGA	GGGATCCC		GGGATCCC	G	GCAGGGCT
2556	GATGGAGC	GGCTAGCTACAACGA	CCTGCGGG		CCCGCAGG	G	GCTCCATC
2561	GAGAGGAT	GGCTAGCTACAACGA	GGAGCCCT		AGGGCTCC	A	ATCCTCTC
2570	AGCAGCGT	GGCTAGCTACAACGA	GGAGAGGA		TCCTCTCC	A	ACGCTGCT
2572	AGAGCAGC	GGCTAGCTACAACGA	GTGGAGAG		CTCTCCAC	G	GCTGCTCT
2575	TGCAGAGC	GGCTAGCTACAACGA	AGCGTGGA		TCCAGGCT	G	GCTCTGCA
2580	CAGGCTGC	GGCTAGCTACAACGA	AGAGCAGC		GCTGCTCT	G	GCAGCCTG
2583	GCACAGGC	GGCTAGCTACAACGA	TGCAGAGC		GCTCTGCA	G	GCCTGTGC
2587	CGTAGCAC	GGCTAGCTACAACGA	AGGCTGCA		TGCAGCCT	G	GTGCTACG
2589	GCCGTAGC	GGCTAGCTACAACGA	ACAGGCTG		CAGCCTGT	G	GCTACGGC
2592	GTGCGCGT	GGCTAGCTACAACGA	AGCACAGG		CCTGTGCT	A	ACGGCGAC
2595	CATGTCGC	GGCTAGCTACAACGA	CGTAGCAC		GTGCTACG	G	GCGACATG
2598	CTCCATGT	GGCTAGCTACAACGA	CGCCGTAG		CTACGGCG	A	ACATGGAG
2600	TTCTCCAT	GGCTAGCTACAACGA	GTCCCGGT		ACGSCGAC	A	ATGGAGAA
2607	CAGCTTGT	GGCTAGCTACAACGA	TCTCCATG		CATGAGA	A	ACAAGCTG
2611	CAACACAGC	GGCTAGCTACAACGA	TGTCTCTC		GAGAACAA	G	GCTGTTTG
2614	CCGCAAAAC	GGCTAGCTACAACGA	AGCTTGTT		AACAAGCT	G	GTTTGGCG
2618	ATCCCCGC	GGCTAGCTACAACGA	AAACAGCT		AGCTGTTT	G	GCGGGGAT
2624	CGCCGAAT	GGCTAGCTACAACGA	CCCCGCCAA		TTGCGGGG	A	ATTCGGCG
2629	CGTCCCGC	GGCTAGCTACAACGA	CGAATCCC		GGGATTCG	G	GCGGGACG
2634	CAGCCCGT	GGCTAGCTACAACGA	CCCGCCGA		TCGCGGGG	A	ACGGGCTG
2638	GGAGCAGC	GGCTAGCTACAACGA	CGTCCCG		CGGGACGG	G	GCTGCTCC
2641	GCAGGAGC	GGCTAGCTACAACGA	AGCCCGTC		GACGGGCT	G	GCTCCTGC
2647	CCAAACGC	GGCTAGCTACAACGA	AGGAGCAG		CTGCTCCT	G	GCCTTTGG
2649	CACCAAAC	GGCTAGCTACAACGA	GCAGGAGC		GCTCCTGC	G	GTTTGGTG

Table 16

2654	TCATCCAC	GGCTAGCTACAACGA	CAAAACGCA			TGCGTTTG	G	GTGGATGA
2658	GAATCAT	GGCTAGCTACAACGA	CCACAAA			TTTGTGG	A	ATGATTTT
2661	CAAGAAAT	GGCTAGCTACAACGA	CATCCACC			GGTGGATG	A	ATTCTTTG
2668	TCACCAAC	GGCTAGCTACAACGA	AAGAAATC			GATTTCTT	G	GTGGTGA
2672	GGTGTAC	GGCTAGCTACAACGA	CAACAAGA			TCTTGTG	G	GTGACACC
2675	TGAGGTGT	GGCTAGCTACAACGA	CACAACA			TGTTGGTG	A	ACACCTCA
2677	GGTGAGGT	GGCTAGCTACAACGA	GTCAACAA			TTGGTGAC	A	ACCTCACC
2682	GGTGAGGT	GGCTAGCTACAACGA	GAGGTGTC			GACACCTC	A	ACCTCACC
2687	GGTGGGT	GGCTAGCTACAACGA	GAGGTGAG			CTCACCTC	A	ACCCACGC
2691	TTTCGGGT	GGCTAGCTACAACGA	GGGTGAGG			CCTCACCC	A	ACGCGAAA
2693	GTTCGCG	GGCTAGCTACAACGA	GTGGGTGA			TCACCCAC	G	GCGAAAAC
2699	AGGAAGGT	GGCTAGCTACAACGA	TTTCGCGT			ACGCGAAA	A	ACCTTCCT
2711	ACCAGGGT	GGCTAGCTACAACGA	CCTGAGGA			TCCTCAGG	A	ACCTTGGT
2717	CCTCGGAC	GGCTAGCTACAACGA	CAGGTGCC			GGACCTTG	G	GTCCGAGG
2724	AGGACAC	GGCTAGCTACAACGA	CTCGGACC			GGTCCGAG	G	GTGTCCCT
2726	TCAGGGAC	GGCTAGCTACAACGA	ACCTCGGA			TCCGAGGT	G	GTCCCTGA
2734	AGCCATAC	GGCTAGCTACAACGA	TCAGGGAC			GTCCCTGA	G	GTATGGCT
2736	GCAGCCAT	GGCTAGCTACAACGA	ACTCAGGG			CCCTGAGT	A	ATGGCTGC
2739	CACGCAGC	GGCTAGCTACAACGA	CATACTCA			TGAGTATG	G	GCTGCGTG
2742	CACCACGC	GGCTAGCTACAACGA	AGCCATAC			GTATGGGT	G	GGTGGTG
2744	TTACCCAC	GGCTAGCTACAACGA	GCAGCCAT			ATGGCTGC	G	GTGGTGAA
2747	AAGTTTAC	GGCTAGCTACAACGA	CACGCAGC			GCTGGTG	G	GTGAACTT
2751	CCGCAAGT	GGCTAGCTACAACGA	TCACCACG			CGTGGTGA	A	ACTTGCGG
2755	TCTTCGCG	GGCTAGCTACAACGA	AAGTTTAC			GTGAACTT	G	GCGGAAGA
2762	ACCACTGT	GGCTAGCTACAACGA	CTTCGCGA			TGCGGAAG	A	ACAGTGGT
2765	TTACCCAC	GGCTAGCTACAACGA	TGTCTTCC			GGAAGACA	G	GTGGTGAA
2768	AAGTTTAC	GGCTAGCTACAACGA	CACCTGCT			AGACAGTG	G	GTGAACTT
2772	AGGGAAGT	GGCTAGCTACAACGA	TCACCAC			AGTGGTGA	A	ACTTCCCT
2780	TCTTCTAC	GGCTAGCTACAACGA	AGGGAAGT			ACTTCCCT	G	GTAGAAGA
2787	GGCCTCGT	GGCTAGCTACAACGA	CTTCTACA			TGTAGAAG	A	ACGAGGCC
2792	CCCAGGGC	GGCTAGCTACAACGA	CTCGTCTT			AAGACGAG	G	GCCTGGG

Table 16

2799	CGTGGCAC	GGCTAGCTACAACGA	CCAGGGCC		GGCCCTGG G GTGGCAG
2802	AGCCGTGC	GGCTAGCTACAACGA	CACCAGG		CCTGGGTG G GCAGGGCT
2804	AAAGCCGT	GGCTAGCTACAACGA	GCCACCA		TGGGTGGC A ACGCTTTT
2807	ACAAAAGC	GGCTAGCTACAACGA	CGTGCCAC		GTGGCAGC G GCTTTTGT
2813	ATCTGAAC	GGCTAGCTACAACGA	AAAAGCCG		CGGCTTTT G GTTCAGAT
2819	GCCGGCAT	GGCTAGCTACAACGA	CTGAACAA		TTGTTTCAG A ATGCCGGC
2821	GGCCCGGC	GGCTAGCTACAACGA	ATCTGAAC		GTTCAGAT G GCCGGCCC
2825	CCGTGGGC	GGCTAGCTACAACGA	CGGCATCT		AGATGCCG G GCCACAGG
2829	TAGGCGGT	GGCTAGCTACAACGA	GGGCCGGC		GCCGGCCC A ACGGCCTA
2832	GAATAGGC	GGCTAGCTACAACGA	CGTGGGCC		GGCCACAG G GCCTATTTC
2836	AGGGGAAT	GGCTAGCTACAACGA	AGGCCGTG		CAGGGCCT A ATTCCCTT
2845	GGCCGCAC	GGCTAGCTACAACGA	CAGGGGAA		TTCCCTTG G GTGGGGCC
2847	CAGGCCGC	GGCTAGCTACAACGA	ACCAGGGG		CCCCTGGT G GCGGCCTG
2850	CAGCAGGC	GGCTAGCTACAACGA	CGCACCCAG		CTGGTGGC G GCCTGCTG
2854	CCAGCAGC	GGCTAGCTACAACGA	AGGCCGCA		TGCGGCCCT G GCTGCTGG
2857	TATCCAGC	GGCTAGCTACAACGA	AGCAGGCC		GGCTGTGT G GCTGGATA
2862	CCGGGTAT	GGCTAGCTACAACGA	CCAGCAGC		GCTGTGG A ATACCCGG
2864	GTCCGGGT	GGCTAGCTACAACGA	ATCCAGCA		TGCTGGAT A ACCGGGAC
2870	TCCAGGGT	GGCTAGCTACAACGA	CCGGGTAT		ATACCCGG A ACCCTGGA
2879	CTCTGCAC	GGCTAGCTACAACGA	CTCCAGGG		CCCTGGAG G GTGCAGAG
2881	CGCTCTGC	GGCTAGCTACAACGA	ACCTCCAG		CTGGAGGT G GCAGAGCG
2886	GTAGTCGC	GGCTAGCTACAACGA	TCTGCACC		GGTGCAGA G GCGACTAC
2889	GGAGTAGT	GGCTAGCTACAACGA	CGCTCTGC		GCAGAGCG A ACTACTCC
2892	GCTGGAGT	GGCTAGCTACAACGA	AGTCGCTC		GAGCGACT A ACTCCAGC
2898	GGCATAGC	GGCTAGCTACAACGA	TGGAGTAG		CTACTCCA G GCTATGCC
2901	COGGGCAT	GGCTAGCTACAACGA	AGCTGGAG		CTCCAGCT A ATGCCCGG
2903	GTCCGGGC	GGCTAGCTACAACGA	ATAGCTGG		CCAGCTAT G GCCGGGAC
2909	ATGGAGGT	GGCTAGCTACAACGA	COGGGCAT		ATGCCCGG A ACCTCCAT
2915	GCTCTGAT	GGCTAGCTACAACGA	GGAGGTCC		GGACCTCC A ATCAGAGC
2921	AGACTGGC	GGCTAGCTACAACGA	TCTGATGG		CCATCAGA G GCGAGTCT
2925	GGTGAGAC	GGCTAGCTACAACGA	TGGCTCTG		CAGAGCCA G GTCTCACC

Table 16

2930	TTGAAGGT	GGCTAGCTACAACGA	GAGACTGG		CCAGTCTC A ACCTTCAA	
2937	GCCGCGGT	GGCTAGCTACAACGA	TGAAGGTG		CACCTTCA A ACCGCGGC	
2940	GAAGCCGC	GGCTAGCTACAACGA	GTTGAAG		CTTCAACC G GGGCTTTC	
2943	CTTGAAGC	GGCTAGCTACAACGA	CGCGGTTG		CAACCGCG G GCTTCAAG	
2951	CTCCAGC	GGCTAGCTACAACGA	CTTGAAGC		GCTTCAAG G GCTGGGAG	
2961	ACGCATGT	GGCTAGCTACAACGA	TCCTCCCA		TGGGAGGA A ACATGCGT	
2963	CGACGCAT	GGCTAGCTACAACGA	GTTCTCTC		GGAGGAAC A ATGCGTCG	
2965	TGCACGCG	GGCTAGCTACAACGA	ATGTTCTT		AGGAACAT G GCGTCGCA	
2967	TTTGCAC	GGCTAGCTACAACGA	GCATGTTT		GAACATGC G GTCGCAAA	
2970	GAGTTTGC	GGCTAGCTACAACGA	GACGCATG		CATGCGTC G GCAAACTC	
2974	CAAGAGAT	GGCTAGCTACAACGA	TTGGGACG		CGTCGCAA A ACTCTTTG	
2984	CGCAAGAC	GGCTAGCTACAACGA	CCCAAGA		TCTTTGGG G GTCTTGCG	
2989	TCAGCCGC	GGCTAGCTACAACGA	AGACCCCT		GGGGTCTT G GCGGCTGA	
2992	ACTTCAGC	GGCTAGCTACAACGA	CGCAAGAC		GTCTTGCG G GCTGAAGT	
2998	TGTGACAC	GGCTAGCTACAACGA	TTGAGCCG		CGGCTGAA G GTGTCACA	
3000	GCTGTGAC	GGCTAGCTACAACGA	ACTTCAGC		GCTGAAGT G GTCACAGC	
3003	CAGGCTGT	GGCTAGCTACAACGA	GACACTTC		GAAGTGC A ACAGCCTG	
3006	AAACAGGC	GGCTAGCTACAACGA	TGTGACAC		GTGTCACA G GCTGTTT	
3010	CCAGAAAC	GGCTAGCTACAACGA	AGGCTGTG		CACAGCCT G GTTCTGCG	
3018	CTGCAAAAT	GGCTAGCTACAACGA	CCAGAAAC		GTTTCTGG A ATTTGCAG	
3022	TCACCTGC	GGCTAGCTACAACGA	AAATCCAG		CTGGATTT G GCAGGTGA	
3026	CTGTTTAC	GGCTAGCTACAACGA	CTGCAAAAT		ATTTGCAG G GTGAACAG	
3030	GAGGCTGT	GGCTAGCTACAACGA	TCACCTGC		GCAGGTGA A ACAGCCTC	
3033	CTGGAGGC	GGCTAGCTACAACGA	TGTTCAAC		GGTGAACA G GCCTCCAG	
3041	CACACCGT	GGCTAGCTACAACGA	CTGGAGGC		GCCTCCAG A ACGGTGTG	
3044	GTGCACAC	GGCTAGCTACAACGA	CGTCTGGA		TCCAGACG G GTGTGCAC	
3046	TGGTGCAC	GGCTAGCTACAACGA	ACCGTCTG		CAGACGGT G GTGCACCA	
3048	GTTGGTGC	GGCTAGCTACAACGA	ACACCGTC		GACGGTGT G GCACCAAC	
3050	ATGTTGGT	GGCTAGCTACAACGA	GCACACCG		CGGTGTGC A ACCAACAT	
3054	GTAGATGT	GGCTAGCTACAACGA	TGGTGCAC		GTGCACCA A ACATCTAC	
3056	TTGTAGAT	GGCTAGCTACAACGA	GTTGGTGC		GCACCAAC A ATCTACAA	

Table 16

3060	GATCTTGT	GGCTAGCTACAACGA	AGATGTTG	CAACATCT A ACAAGATC
3065	AGGAGGAT	GGCTAGCTACAACGA	CTTGAGAA	TCTACAAG A ATCCTCCT
3073	CCTGCAGC	GGCTAGCTACAACGA	AGGAGGAT	ATCCTCCT G GCTGCAGG
3076	ACGCTGTC	GGCTAGCTACAACGA	AGCAGGAG	CTCCTGCT G GCAGGCCGT
3080	CTGTACGC	GGCTAGCTACAACGA	CTGCAGCA	TGCTGCAG G GCGTACAG
3082	ACCTGTAC	GGCTAGCTACAACGA	GCCTGCAG	CTGCAGGC G GTACAGGT
3084	AAACTGT	GGCTAGCTACAACGA	ACGCTGTC	GCAGGCCGT A ACAGGTTT
3088	CGTGAAC	GGCTAGCTACAACGA	CTGTACGC	GCGTACAG G GTTTCACG
3093	ACATGCGT	GGCTAGCTACAACGA	GAACCTGT	CAGGTTTC A ACGCATGT
3095	ACACATGC	GGCTAGCTACAACGA	GTGAACCC	GGTTCTAC G GCATGTGT
3097	GCACACAT	GGCTAGCTACAACGA	GTGAACAA	TTCACGC A ATGTGTGC
3099	CAGCACAC	GGCTAGCTACAACGA	ATGCGTGA	TCACGCAT G GTGTGCTG
3101	TGCAGCAC	GGCTAGCTACAACGA	ACATGCGT	ACGCATGT G GTGCTGCA
3103	GCTGCAGC	GGCTAGCTACAACGA	ACACATGC	GCATGTGT G GCTGCAGC
3106	GGAGCTGC	GGCTAGCTACAACGA	AGCACACA	TGTGTGCT G GCAGCTCC
3109	ATGGAGC	GGCTAGCTACAACGA	TGCAGCAC	GTGCTGCA G GCTCCCAT
3115	GATGAAAT	GGCTAGCTACAACGA	GGGAGCTG	CAGTCCC A ATTCATC
3120	TTGCTGAT	GGCTAGCTACAACGA	GAAATGGG	CCCATTTT A ATCAGCAA
3124	AAACTTGC	GGCTAGCTACAACGA	TGATGAAA	TTTCATCA G GCAAGTTT
3128	TTCCAAAC	GGCTAGCTACAACGA	TTGCTGAT	ATCAGCAA G GTTGGAA
3138	TGTGGGT	GGCTAGCTACAACGA	TCTTCCAA	TTGGAAGA A ACCCCACA
3143	AAAAATGT	GGCTAGCTACAACGA	GGGTTCT	AGAACCCC A ACATTTT
3145	GGAAAAAT	GGCTAGCTACAACGA	GTGGGGTT	AACCCAC A ATTTTCC
3154	TGACGGC	GGCTAGCTACAACGA	AGGAAAA	TTTTTCT G GCGGTCA
3156	GATGACG	GGCTAGCTACAACGA	GCAGGAAA	TTTCTGCG G GCGTCATC
3158	GAGATGAC	GGCTAGCTACAACGA	GCAGGAGA	TCCTGCGC G GTCATCTC
3161	TCAGAGAT	GGCTAGCTACAACGA	GACGCGCA	TGCGCGTC A ATCTCTGA
3168	GGCCGTGT	GGCTAGCTACAACGA	CAGAGATG	CATCTCTG A ACAGGGCC
3170	GAGGCGGT	GGCTAGCTACAACGA	GTCAGAGA	TCTCTGAC A ACGGCTCT
3173	AGGGAGGC	GGCTAGCTACAACGA	CGTGTGAG	CTGACACG G GCTCCCT
3183	GGAGTAGC	GGCTAGCTACAACGA	AGAGGGAG	CTCCCTCT G GCTACTCC

Table 16

3186	GATGGAGT	GGCTAGCTACAACGA	AGCAGAGG			CCTCTGCT A	ACTCCATC
3191	TTCAGGAT	GGCTAGCTACAACGA	GGAGTAGC			GCTACTCC A	ATCCTGAA
3200	TTCTTGGC	GGCTAGCTACAACGA	TTTCAGGA			TCCTGAAA G	GCCAAGAA
3207	CCCTGCGT	GGCTAGCTACAACGA	TCTTGGCT			AGCCAAGA A	ACGCAGGG
3209	ATCCCTGC	GGCTAGCTACAACGA	GTTCTTGG			CCAAGAAC G	GCAGGGAT
3215	AGCGACAT	GGCTAGCTACAACGA	CCCTGCGT			ACGCAGGG A	ATGTCGCT
3217	CCAGCGAC	GGCTAGCTACAACGA	ATCCCTGC			GCAGGAT G	GTCGCTGG
3220	CCCCCAGC	GGCTAGCTACAACGA	GACATCCC			GGGATGTC G	GCTGGGGG
3227	CCCTTGGC	GGCTAGCTACAACGA	CCCCAGCG			CGCTGGGG G	GCCAAGGG
3234	GGCGGCGC	GGCTAGCTACAACGA	CCTTGGCC			GGCCAAGG G	GCGCCGCC
3236	CCGGCGGC	GGCTAGCTACAACGA	GCCTTTGG			CCAAGGGC G	GCCGCCGG
3239	GGGCCGGC	GGCTAGCTACAACGA	GGCGCCCT			AGGGCGGC G	GCCGGCCC
3243	CAGAGGGC	GGCTAGCTACAACGA	CGGGGGCG			CGCGCGCG G	GCCCTCTG
3250	CGGAGGGC	GGCTAGCTACAACGA	AGAGGGCC			GGCCCTCT G	GCCCTCCG
3260	TGCACGGC	GGCTAGCTACAACGA	CTCGGAGG			CCTCCGAG G	GCCTGCA
3263	CACTGCAC	GGCTAGCTACAACGA	GGCCTCGG			CCGAGGCC G	GTGCAGTG
3265	GCCACTGC	GGCTAGCTACAACGA	ACGGCCTC			GAGGCCGT G	GCAGTGGC
3268	ACAGCCAC	GGCTAGCTACAACGA	TGCACGGC			GCCGTGCA G	GTGGCTGT
3271	GGCACAGC	GGCTAGCTACAACGA	CACTGCAC			GTGCAGTG G	GCTGTGCC
3274	GGTGGCAC	GGCTAGCTACAACGA	AGCCACTG			CAGTGGCT G	GTGCCACC
3276	TTGGTGGC	GGCTAGCTACAACGA	ACAGCCAC			GTGGCTGT G	GCCACCAC
3279	TGCTTGGT	GGCTAGCTACAACGA	GGCACAGC			GCTGTGCC A	ACCAAGCA
3284	AGGAATGC	GGCTAGCTACAACGA	TTGGTGGC			GCCACCAA G	GCATTCCT
3286	GCAGGAAT	GGCTAGCTACAACGA	GCTTGGTG			CACCAAGC A	ATTCTCTG
3292	GCTTGAGC	GGCTAGCTACAACGA	AGGAATGC			GCATTCCT G	GCTCAAGC
3298	GAGTCAGC	GGCTAGCTACAACGA	TTGAGCAG			CTGCTCAA G	GCTGACTC
3302	TGTCGAGT	GGCTAGCTACAACGA	CAGCTTGA			TCAAGCTG A	ACTCGACA
3307	CACGGTGT	GGCTAGCTACAACGA	CGAGTCAG			CTGACTCG A	ACACCGTG
3309	GACACGGT	GGCTAGCTACAACGA	GTCGAGTC			GACTCGAC A	ACCGTGTC
3312	GGTGACAC	GGCTAGCTACAACGA	GGTGTCGA			TCGACACC G	GTGTCACC
3314	TAGGTGAC	GGCTAGCTACAACGA	ACGGTGTG			GACACCGT G	GTACACCTA

Table 16

3317	ACGTAGGT	GGCTAGCTACAACGA	GACACGGT		ACCGTGTG A	ACCTACGT
3321	TGGCACGT	GGCTAGCTACAACGA	AGGTGACA		TGTCACT A	ACGTGCCA
3323	AGTGGCAC	GGCTAGCTACAACGA	GTAGGTGA		TCACCTAC G	GTGCCACT
3325	GGAGTGGC	GGCTAGCTACAACGA	ACGTAGGT		ACCTACGT G	GCCACTCC
3328	CCAGGAGT	GGCTAGCTACAACGA	GGCAGGTA		TACGTGCC A	ACTCCTGG
3337	TGAGTGAC	GGCTAGCTACAACGA	CCCAGGAG		CTCCTGGG G	GTCACCTA
3340	TCCTGAGT	GGCTAGCTACAACGA	GACCCAG		CTGGGGTC A	ACTCAGGA
3347	TGGGCTGT	GGCTAGCTACAACGA	CCTGAGTG		CACTCAGG A	ACAGCCCA
3350	GTCTGGGC	GGCTAGCTACAACGA	TGTCCTGA		TCAGGACA G	GCCCAGAC
3356	AGCTGGCT	GGCTAGCTACAACGA	CTGGGCTG		CAGCCCAG A	ACGCAGCT
3358	TCAGCTGC	GGCTAGCTACAACGA	GTCTGGGC		GCCCAGAC G	GCAGCTGA
3361	GACTCAGC	GGCTAGCTACAACGA	TGGTCTG		CAGACGCA G	GCTGAGTC
3366	CTTCCGAC	GGCTAGCTACAACGA	TCAGCTGC		GCAGCTGA G	GTCGGAAG
3373	CCGGGAGC	GGCTAGCTACAACGA	TTCGACT		AGTCGGAA G	GCTCCCGG
3383	AGCGTCGT	GGCTAGCTACAACGA	CCCAGGGA		TCCCGGGG A	ACGACGCT
3386	GTACGCGT	GGCTAGCTACAACGA	CGTCCCGG		CGGGGACG A	ACGCTGAC
3388	CAGTCAGC	GGCTAGCTACAACGA	GTCTGCC		GGGACGAC G	GCTGACTG
3392	AGGCGAGT	GGCTAGCTACAACGA	CAGCGTCG		CGACGCTG A	ACTGCCCT
3395	TCCAGGGC	GGCTAGCTACAACGA	AGTCAGCG		CGCTGACT G	GCCTGGA
3404	GCTGGCGC	GGCTAGCTACAACGA	CTCCAGGG		CCCTGGAG G	GCCGACGC
3407	TTGGCTGC	GGCTAGCTACAACGA	GGCTCCA		TGGAGGCC G	GCAGCCAA
3410	GGGTTGGC	GGCTAGCTACAACGA	TGGGCTCT		AGGCCGCA G	GCCAACCC
3414	TGCCGGGT	GGCTAGCTACAACGA	TGGCTGCG		CGCAGCCA A	ACCCGGCA
3419	GGCAGTGC	GGCTAGCTACAACGA	CGGGTTGG		CCAACCCG G	GCACTGCC
3421	AGGCGAGT	GGCTAGCTACAACGA	GCCGGGTT		AACCCGGC A	ACTGCCCT
3424	CTGAGGGC	GGCTAGCTACAACGA	AGTGCCGG		CCGGCACT G	GCCTCAG
3432	CTTGAAGT	GGCTAGCTACAACGA	CTGAGGGC		GCCCTCAG A	ACTTCAAG
3440	AGGATGGT	GGCTAGCTACAACGA	CTTGAAGT		ACTTCAAG A	ACCATCCT
3443	TCCAGGAT	GGCTAGCTACAACGA	GGTCTTGA		TCAAGACC A	ATCCTGGA
3450	CCATCAGT	GGCTAGCTACAACGA	CCAGGATG		CATCCTGG A	ACTGATGG
3454	GTGGCCAT	GGCTAGCTACAACGA	CAGTCCAG		CTGGACTG A	ATGGCCAC

Table 16

3457	CGGGTGGC	GGCTAGCTACAAACGA	CATCAGTC	GACTGATG	G	GCCACCCG
3460	GGGCGGGT	GGCTAGCTACAAACGA	GGCCATCA	TGATGGC	A	ACCCGCC
3464	CTGTGGC	GGCTAGCTACAAACGA	GGGTGGC	GGCCACCC	G	GCCACACG
3468	CTGGCTGT	GGCTAGCTACAAACGA	GGCGGGT	ACCCGCC	A	ACAGCCAG
3471	GGCCTGGC	GGCTAGCTACAAACGA	TGTGGCG	CGCCACA	G	GCCAGGCC
3476	CTCTCGGC	GGCTAGCTACAAACGA	CTGGCTGT	ACAGCCAG	G	GCCGAGAG
3483	GTGTCTGC	GGCTAGCTACAAACGA	TCTCGGCC	GGCCGAGA	G	GCAGACAC
3487	GCTGTGT	GGCTAGCTACAAACGA	CTGCTCTC	GAGAGCAG	A	ACACCAGC
3489	CTGCTGGT	GGCTAGCTACAAACGA	GTCTGCTC	GAGCAGAC	A	ACCAGCAG
3493	AGGGCTGC	GGCTAGCTACAAACGA	TGGTGTCT	AGACACCA	G	GCAGCCCT
3496	GACAGGGC	GGCTAGCTACAAACGA	TGCTGGTG	CACCAGCA	G	GCCTGTGC
3501	GGCGTGAC	GGCTAGCTACAAACGA	AGGGCTGC	GCAGCCCT	G	GTACACGC
3504	CCCGGGT	GGCTAGCTACAAACGA	GACAGGGC	GCCTGTG	A	ACGCCGGG
3506	AGCCCGGC	GGCTAGCTACAAACGA	GTGACAGG	CTGTCTAC	G	GCAGGGCT
3511	CGTAGAGC	GGCTAGCTACAAACGA	CCGGCTGT	CAGCCCG	G	GCTCTACG
3516	TGGGACGT	GGCTAGCTACAAACGA	AGAGCCG	CGGGCTCT	A	ACGTCCCA
3518	CCTGGAC	GGCTAGCTACAAACGA	GTAGAGCC	GGCTCTAC	G	GTCCCAGG
3535	TGGCCGC	GGCTAGCTACAAACGA	CCCTCCCT	AGGGAGGG	G	GCAGGCCA
3538	GTGTGGC	GGCTAGCTACAAACGA	CGCCCTC	GAGGGGCG	G	GCCACAC
3542	CTGGGTGT	GGCTAGCTACAAACGA	GGGCCGCC	GGCGGCC	A	ACACCCAG
3544	GCCTGGGT	GGCTAGCTACAAACGA	GTGGCCG	CGGCCAC	A	ACCCAGGC
3550	GTGCGGC	GGCTAGCTACAAACGA	CTGGGTGT	ACACCCAG	G	GCAGGCAC
3554	AGCGGTGC	GGCTAGCTACAAACGA	GGGCCCTGG	CGAGGCC	G	GCAGCGCT
3556	CCAGCGGT	GGCTAGCTACAAACGA	GCGGGCT	AGCCCGC	A	ACCGCTGG
3559	CTCCAGC	GGCTAGCTACAAACGA	GGTGCGG	CCCGACC	G	GCTGGGAG
3566	CCTCAGAC	GGCTAGCTACAAACGA	TCCCAGG	CGCTGGGA	G	GTCTGAGG
3573	ACTCAGGC	GGCTAGCTACAAACGA	CTCAGACT	AGTCTGAG	G	GCCTGAGT
3579	ACACTCAC	GGCTAGCTACAAACGA	TCAGGCCT	AGGCCTGA	G	GTGAGTGT
3583	CCAAACAC	GGCTAGCTACAAACGA	TCACTCAG	CTGAGTGA	G	GTGTTGG
3585	GGCCAAAC	GGCTAGCTACAAACGA	ACTCACTC	GAGTGA	G	GTGTTGCC
3590	GCCTCGGC	GGCTAGCTACAAACGA	CAACACT	AGTGTG	G	GCCGAGGC

Table 16

3596	ATGCAGGC	GGCTAGCTACAACGA	CTCGGCCA		TGCCCCGAG G GCCTGCAT
3600	GGACATGC	GGCTAGCTACAACGA	AGGCTCG		CGAGGCCT G GCATGTCC
3602	CCGGACAT	GGCTAGCTACAACGA	GCAGCCT		AGGCCTGC A ATGTCCGG
3604	AGCCGGAC	GGCTAGCTACAACGA	ATGCAGGC		GCCTGCAT G GTCCGGCT
3609	CCTTCAGC	GGCTAGCTACAACGA	CGGACATG		CATGTCCG G GCTGAAGG
3616	CACTCAGC	GGCTAGCTACAACGA	CTTCAGCC		GGCTGAAG G GCTGAGTG
3621	CCGGACAC	GGCTAGCTACAACGA	TCAGCCTT		AAGGCTGA G GTGTCCGG
3623	AGCCGGAC	GGCTAGCTACAACGA	ACTCAGCC		GGTGAGT G GTCCGGCT
3628	GCCTCAGC	GGCTAGCTACAACGA	CGGACACT		AGTGTCCG G GCTGAGGC
3634	GCTCAGGC	GGCTAGCTACAACGA	CTCAGCCG		CGGCTGAG G GCCTGAGC
3640	ACACTCGC	GGCTAGCTACAACGA	TCAGGCCT		AGGCCTGA G GCGAGTGT
3644	CTGGACAC	GGCTAGCTACAACGA	TCGCTCAG		CTGAGCGA G GTGTCCAG
3646	GGCTGGAC	GGCTAGCTACAACGA	ACTCGCTC		GAGCGAGT G GTCCAGCC
3651	CCCTTGGC	GGCTAGCTACAACGA	TGGACACT		AGTGTCCA G GCCAAGGG
3658	CACTCAGC	GGCTAGCTACAACGA	CCTTGGCT		AGCCAAGG G GCTGAGTG
3663	CTGGACAC	GGCTAGCTACAACGA	TCAGCCCT		AGGCTGA G GTGTCCAG
3665	TGCTGGAC	GGCTAGCTACAACGA	ACTCAGCC		GGTGAGT G GTCCAGCA
3670	AGGTGTGC	GGCTAGCTACAACGA	TGGACACT		AGTGTCCA G GCACACCT
3672	GCAGGTGT	GGCTAGCTACAACGA	GCTGGACA		TGTCAGC A ACACCTGC
3674	CGGCAGGT	GGCTAGCTACAACGA	GTGCTGGA		TCCAGCAC A ACCTGCCG
3678	AAGACGGC	GGCTAGCTACAACGA	AGGTGTGC		GCACACCT G GCGTCTT
3681	GTGAAGAC	GGCTAGCTACAACGA	GGCAGGTG		CACCTGCC G GTCTTCAC
3687	GCGGAAGT	GGCTAGCTACAACGA	GAAGACGG		CCGTCTT A ACTTCCC
3695	CAGCCTGT	GGCTAGCTACAACGA	GCGGNAAGT		ACTTCCCC A ACAGGCTG
3699	GCGCCAGC	GGCTAGCTACAACGA	CTGTGGGG		CCCCACAG G GCTGGGCG
3703	CCGAGCGC	GGCTAGCTACAACGA	CAGCCTGT		ACAGGCTG G GCGCTCGG
3705	AGCCGAGC	GGCTAGCTACAACGA	GCAGCCT		AGGTGGC G GCTCGGCT
3710	GGTGGAGC	GGCTAGCTACAACGA	CGAGCGCC		GGCGTCCG G GCTCCACC
3715	CCTGGGGT	GGCTAGCTACAACGA	GGAGCCGA		TCGGCTCC A ACCCCAGG
3723	AAGCTGGC	GGCTAGCTACAACGA	CCTGGGGT		ACCCACAGG G GCCAGCTT
3727	GGAAAGC	GGCTAGCTACAACGA	TGGCCCTG		CAGGGCCA G GCTTTTCC

Table 16

3737	CTCCTGGT	GGCTAGCTACAACGA	GAGGAAAA		TTTTCTC A ACCAGGAG
3744	AGCCGGGC	GGCTAGCTACAACGA	TCCTGGTG		CACAGGA G GCCGGCT
3749	GTGGAAGC	GGCTAGCTACAACGA	CGGGCTCC		GGAGCCG G GCTTCAC
3755	TGGGGAGT	GGCTAGCTACAACGA	GGAAGCCG		CGGCTCC A ACTCCCA
3762	TCCTATGT	GGCTAGCTACAACGA	GGGGAGTG		CACCCCC A ACATAGGA
3764	ATTCCTAT	GGCTAGCTACAACGA	GTGGGGAG		CTCCCCA A ATAGGAAT
3770	TGGACTAT	GGCTAGCTACAACGA	TCCTATGT		ACATAGGA A ATAGTCCA
3773	GGATGGAC	GGCTAGCTACAACGA	TATTCCTA		TAGGAATA G GTCCATCC
3777	CTGGGGAT	GGCTAGCTACAACGA	GGAATAAT		AATAGTCC A ATCCCOAG
3785	TGGCGAAT	GGCTAGCTACAACGA	CTGGGGAT		ATCCCOAG A ATTCGCCA
3789	ACAATGGC	GGCTAGCTACAACGA	GAATCTGG		CCAGATTC G GCCATTGT
3792	TGAACAAT	GGCTAGCTACAACGA	GCGGAATC		GAATGCC A ATTGTTCA
3795	GGGTGAAC	GGCTAGCTACAACGA	AATGGCGA		TCGCCATT G GTTCACCC
3799	CGAGGGGT	GGCTAGCTACAACGA	GAACAATG		CATTGTTT A ACCCTCG
3806	GGCAGGGC	GGCTAGCTACAACGA	GAGGGGTG		CACCCCTC G GCCCTGCC
3811	AGGAGGGC	GGCTAGCTACAACGA	AGGGCGAG		CTCGCCCT G GCCCTCCT
3821	TGGAAGGC	GGCTAGCTACAACGA	AAAGGAGG		CCTCCTTT G GCCTTCCA
3828	GTGGGGGT	GGCTAGCTACAACGA	GGAAGGCA		TGCCTTCC A ACCCCAC
3834	TGGATGGT	GGCTAGCTACAACGA	GGGGGTGG		CCACCCCC A ACCATCCA
3837	ACCTGGAT	GGCTAGCTACAACGA	GGTGGGGG		CCCCCACC A ATCCAGGT
3843	GTCTCCAC	GGCTAGCTACAACGA	CTGGATGG		CCATCCAG G GTGGAGAC
3849	CTCAGGGT	GGCTAGCTACAACGA	CTCCACCT		AGGTGGAG A ACCCTGAG
3861	CCCAGGGT	GGCTAGCTACAACGA	CCTTCTCA		TGAGAAGG A ACCCTGGG
3870	CCCAGAGC	GGCTAGCTACAACGA	TCCCAGGG		CCCTGGGA G GCTCTGGG
3879	CTCCAAAT	GGCTAGCTACAACGA	TCCCAGAG		CTCTGGGA A ATTTGGAG
3886	TTGGTCAC	GGCTAGCTACAACGA	TCCAAATT		AATTTGGA G GTGACCAA
3889	CCTTTGGT	GGCTAGCTACAACGA	CACTCCAA		TTGGAGTG A ACCAAAGG
3896	GGGCACAC	GGCTAGCTACAACGA	CTTTGGTC		GACCAAAG G GTGTGCC
3898	CAGGGCAC	GGCTAGCTACAACGA	ACCTTTGG		CCAAAGGT G GTGCCCTG
3900	TACAGGGC	GGCTAGCTACAACGA	ACACCTTT		AAAGGTGT G GCCCTGTA
3905	CTGTGTAC	GGCTAGCTACAACGA	AGGGCACA		TGTGCCCT G GTACACAG

Table 16

3907	GCCTGTGT	GGCTAGCTACAACGA	ACAGGGCA		TGCCCTGT	A	ACACAGGC
3909	TGCCCTGT	GGCTAGCTACAACGA	GTACAGGG		CCCTGTAC	A	ACAGGCGA
3913	GTCTCGC	GGCTAGCTACAACGA	CTGTGTAC		GTACACAG	G	GCAGGAC
3919	TGCAGGGT	GGCTAGCTACAACGA	CCTCGCT		AGCGAGG	A	ACCCTGCA
3924	CCAGGTGC	GGCTAGCTACAACGA	AGGTCCT		AGGACCT	G	GCACCTGG
3926	ATCCAGGT	GGCTAGCTACAACGA	GCAGGTC		GACCCCTG	A	ACCTGGAT
3932	ACCCCCAT	GGCTAGCTACAACGA	CCAGGTGC		GCACCTGG	A	ATGGGGGT
3938	ACAGGGAC	GGCTAGCTACAACGA	CCCCATCC		GGATGGG	G	GTCCCTGT
3944	TGACCCAC	GGCTAGCTACAACGA	AGGGACCC		GGTCCCT	G	GTGGGTCA
3948	AATTGAC	GGCTAGCTACAACGA	CCACAGGG		CCCTGTGG	G	GTCAAAAT
3953	CCCCAAT	GGCTAGCTACAACGA	TTGACCCA		TGGGTCAA	A	ATTGGGGG
3964	CACAGCAC	GGCTAGCTACAACGA	CTCCCCC		GGGGGAG	G	GTGCTGTG
3966	CCACAGC	GGCTAGCTACAACGA	ACCTCCCC		GGGGAGGT	G	GCTGTGGG
3969	ACTCCAC	GGCTAGCTACAACGA	AGCACCTC		GAGGTGCT	G	GTGGGAGT
3975	TATTTTAC	GGCTAGCTACAACGA	TCCCACAG		CTGTGGGA	G	GTAAATA
3980	TTCAGTAT	GGCTAGCTACAACGA	TTTACTCC		GGAGTAA	A	ATACTGAA
3982	TATTCAGT	GGCTAGCTACAACGA	ATTTTACT		AGTAAAT	A	ACTGAATA
3987	TCATATAT	GGCTAGCTACAACGA	TCAGTATT		AAACTGA	A	ATATATGA
3989	ACTCATAT	GGCTAGCTACAACGA	ATTCAGTA		TACTGAAT	A	ATATGAGT
3991	AACTCAT	GGCTAGCTACAACGA	ATATTCAG		CTGAATAT	A	ATGAGTTT
3995	TGAAAAAC	GGCTAGCTACAACGA	TCATATAT		ATATATGA	G	GTTTTC
4003	TTCAAAAC	GGCTAGCTACAACGA	TGAAAAAC		GTTTTC	G	GTTTGA

Seq1 = TERT (Homo sapiens telomerase reverse transcriptase (TERT) mRNA, 4015 bp); Nakamura *et al.*, Science 277 (5328), 955-959 (1997)
 Cut Site = R/Y (Purine/Pyrimidine)

Stem Length = 8 . Core Sequence = GGCTAGCTACAACGA

343
Table 17

Table 17: Anti-TERT HH and G-Cleaver Ribozymes

Alias	Ribozyme Sequence	Length (nt)
HH		
TERT-1051	AGGAGUA CUGAUGAGGCCGUAAGGCCGAA AGGAAGU	36
TERT-1053	UGAGGAG CUGAUGAGGCCGUAAGGCCGAA AGAGGAA	36
TERT-1918	UGAAGCG CUGAUGAGGCCGUAAGGCCGAA AGUCUGG	36
TERT-2383	GAGCCAC CUGAUGAGGCCGUAAGGCCGAA AACUGUC	36
TERT-2485	UGAAGCG CUGAUGAGGCCGUAAGGCCGAA AGGAAGA	36
TERT-2566	GCGUGGA CUGAUGAGGCCGUAAGGCCGAA AGGAUGG	36
TERT-3181	AGUAGCA CUGAUGAGGCCGUAAGGCCGAA AGGGAGG	36
TERT-3691	CUGUGGG CUGAUGAGGCCGUAAGGCCGAA AAGUGAA	36
TERT-3758	AUGUGGG CUGAUGAGGCCGUAAGGCCGAA AGUGGAA	36
TERT-3794	GGUGAAC CUGAUGAGGCCGUAAGGCCGAA AUGGCGA	36
G-Cleaver		
TERT-757	UUGGG UGAUGGCAUGCACUAUGCGCG AACGGCAGAC	36
TERT-2353	UCUGU UGAUGGCAUGCACUAUGCGCG AAGGUAGAGA	36
TERT-3795	GUGAA UGAUGGCAUGCACUAUGCGCG AAUGGCGAAU	36

Table 18

Table 18: Human BACE Hammerhead Ribozyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
9	CCACGCGU C CGCAGCCC	1	GGGUGCG CUGAUGAG X CGAA ACGCGUGG	1776
47	AGCUGGAU U AUGGUGGC	2	GCCACCAU CUGAUGAG X CGAA AUCCAGCU	1777
48	GCUGGAU A UGGUGGCC	3	GGCCACCA CUGAUGAG X CGAA AAUCCAGC	1778
93	GGAGCCCU U GCCCCUGC	4	GCAGGGGC CUGAUGAG X CGAA AGGGCUCC	1779
163	CCGCCCCU C CCAGCCCC	5	GGGGCUGG CUGAUGAG X CGAA AGGGGCGG	1780
221	GCCGAUGU A GCGGGCUC	6	GAGCCCGC CUGAUGAG X CGAA ACAUCGGC	1781
229	AGCGGGCU C CGGAUCCC	7	GGGAUCCG CUGAUGAG X CGAA AGCCCGCU	1782
235	CUCGGGAU C CCAGCCUC	8	GAGGCGUG CUGAUGAG X CGAA AUCCGGAG	1783
243	CCCAGCCU C UCCCCUGC	9	GCAGGGGA CUGAUGAG X CGAA AGGCUGGG	1784
245	CAGCCUCU C CCCUGCUC	10	GAGCAGGG CUGAUGAG X CGAA AGAGGCUG	1785
253	CCCCUGCU C CCGUGCUC	11	GAGCACGG CUGAUGAG X CGAA AGCAGGGG	1786
261	CCCGUGCU C UCGGAUC	12	GAUCCGCA CUGAUGAG X CGAA AGCACGGG	1787
269	CUGCGGAU C UCCCCUGA	13	UCAGGGGA CUGAUGAG X CGAA AUCCGCAG	1788
271	GCGGAUCU C CCCUGACC	14	GGUCAGGG CUGAUGAG X CGAA AGAUCCGC	1789
283	UGACCGCU C UCCACAGC	15	GCUGUGGA CUGAUGAG X CGAA AGCGGUCA	1790
285	ACCGCUCU C CACAGCCC	16	GGGUGUG CUGAUGAG X CGAA AGAGCGGU	1791
334	CCUGGCGU C CUGAUGCC	17	GGCAUCAG CUGAUGAG X CGAA ACGCCAGG	1792
351	CCAAGCU C CCUCUCCU	18	AGGAGAGG CUGAUGAG X CGAA AGCUUGGG	1793
355	AGCUCCCU C UCCUGAGA	19	UCUCAGGA CUGAUGAG X CGAA AGGGAGCU	1794
357	CUCCUCU C CUGAGAAG	20	CUUCUCAG CUGAUGAG X CGAA AGAGGGAG	1795
386	CCCAGACU U GGGGCGAG	21	CUGCCCC CUGAUGAG X CGAA AGUCUGGG	1796
477	CCUGGCU C CUGCUGUG	22	CACAGCAG CUGAUGAG X CGAA AGCCAGGG	1797
531	CACGGCAU C CGGUGCC	23	GGCAGCCG CUGAUGAG X CGAA AUGCCGUG	1798
632	GGGAGCU U UGUGGAGA	24	UCUCCACA CUGAUGAG X CGAA AGCUGCCC	1799
633	GGCAGCUU U GUGGAGAU	25	AUCUCCAC CUGAUGAG X CGAA AAGCUGCC	1800
665	GGGCAAGU C GGGGCGAG	26	CCUGCCCC CUGAUGAG X CGAA ACTUGCCC	1801
677	GCAGGGCU A CUACGUGG	27	CCACGUAG CUGAUGAG X CGAA AGCCCUGC	1802
680	GGGCUACU A CGUGGAGA	28	UCUCCACG CUGAUGAG X CGAA AGUAGCCC	1803
717	CAGACGCU C AACAUCCU	29	AGGAUGUU CUGAUGAG X CGAA AGCGUCUG	1804
723	CUCAACAU C CUGUGGA	30	UCCACGAG CUGAUGAG X CGAA AUGUUGAG	1805
733	UGGUGGAU A CAGGCAGC	31	GCUGCCUG CUGAUGAG X CGAA AUCCACCA	1806
745	GCAGCAGU A ACUUUGCA	32	UGCAAAGU CUGAUGAG X CGAA ACUGCUGC	1807
749	CAGUAACU U UGCAGUGG	33	CCACUGCA CUGAUGAG X CGAA AGUUACUG	1808
750	AGUAACUU U GCAGUGGG	34	CCCACUGC CUGAUGAG X CGAA AAGUUACU	1809
776	CCACCCCU U CCUGCAUC	35	GAUGCAGG CUGAUGAG X CGAA AGGGGUGG	1810
777	CACCCCUU C CUGCAUCG	36	CGAUGCAG CUGAUGAG X CGAA AAGGGGUG	1811
784	UCCUGCAU C GCUACUAC	37	GUAGUAGC CUGAUGAG X CGAA AUGCAGGA	1812
788	GCAUCGCU A CUACCAGA	38	UCUGGUAG CUGAUGAG X CGAA AGCGAUGC	1813
791	UCGCUACU A CCAGAGGC	39	GCCUCUGG CUGAUGAG X CGAA AGUAGCGA	1814
806	GCAGCUGU C CAGCACAU	40	AUGUGCUG CUGAUGAG X CGAA ACAGCUGC	1815
815	CAGCACAU A CCGGACC	41	GGUCCCGG CUGAUGAG X CGAA AUGUGCUG	1816
825	CGGGACCU C CGGAAGGG	42	CCCUUCCG CUGAUGAG X CGAA AGGUCCCG	1817
839	GGGUGUGU A UGUGCCCU	43	AGGGCACA CUGAUGAG X CGAA ACACACCC	1818
848	UGUGCCCU A CACCCAGG	44	CCUGGGUG CUGAUGAG X CGAA AGGGCACA	1819
891	GACCGGU A AGCAUCCC	45	GGGAUGCU CUGAUGAG X CGAA ACCAGGUC	1820
897	GUAAGCAU C CCCAUGG	46	CCAUGGGG CUGAUGAG X CGAA AUGCUUAC	1821
915	CCCAACGU C ACUGUGCG	47	CGCACAGU CUGAUGAG X CGAA ACGUUGGG	1822

Table 18

933	GCCAACAU U GCUGCCAU	48	AUGGCAGC CUGAUGAG X CGAA AUGUUGGC	1823
942	GCUGCCAU C ACUGAAUC	49	GAUUCAGU CUGAUGAG X CGAA AUGGCAGC	1824
950	CACUGAAU C AGACAAGU	50	ACUUGUCU CUGAUGAG X CGAA AUUCAGUG	1825
959	AGACAAGU U CUUCAUCA	51	UGAUGAAG CUGAUGAG X CGAA ACUUGUCU	1826
960	GACAAGUU C UUCAUCA	52	UUGAUGAA CUGAUGAG X CGAA AACUUGUC	1827
962	CAAGUUCU U CAUCAACG	53	CGUUGAUG CUGAUGAG X CGAA AGAACUUG	1828
963	AAGUUCUU C AUCAACGG	54	CCGUUGAU CUGAUGAG X CGAA AAGAACUU	1829
966	UUCUUCAU C AACGGCUC	55	GAGCCGUU CUGAUGAG X CGAA AUGAAGAA	1830
974	CAACGGCU C CAACUGGG	56	CCCAGUUG CUGAUGAG X CGAA AGCCGUUG	1831
990	GAAGGCAU C CUGGGGCU	57	AGCCCCAG CUGAUGAG X CGAA AUGCCUUC	1832
1004	GCUGGCCU A UGUGAGA	58	UCUCAGCA CUGAUGAG X CGAA AGGCCAGC	1833
1014	GCUGAGAU U GCCAGGCC	59	GGCCUGGC CUGAUGAG X CGAA AUCUCAGC	1834
1031	UGACGACU C CCUGGAGC	60	GCUCCAGG CUGAUGAG X CGAA AGUCGUCA	1835
1042	UGGAGCCU U UCUUUGAC	61	GUCAAAGA CUGAUGAG X CGAA AGGCUCCA	1836
1043	GGAGCCUU U CUUUGACU	62	AGUCAAG CUGAUGAG X CGAA AAGGCUCC	1837
1044	GAGCCUUU C UUUGACUC	63	GAGUCAA CUGAUGAG X CGAA AAAGGCUC	1838
1046	GCCUUUCU U UGACUCUC	64	GAGAGUCA CUGAUGAG X CGAA AGAAAGGC	1839
1047	CCUUUCUU U GACUCUCU	65	AGAGAGUC CUGAUGAG X CGAA AAGAAAGG	1840
1052	CUUUGACU C UCUGGUAA	66	UUACCAGA CUGAUGAG X CGAA AGUCAAG	1841
1054	UUGACUCU C UGUUAAAG	67	CUUUAACA CUGAUGAG X CGAA AGAGUCAA	1842
1059	UCUCUGGU A AAGCAGAC	68	GUCUGCUU CUGAUGAG X CGAA ACCAGAGA	1843
1074	ACCCACGU U CCCAACCU	69	AGGUUGGG CUGAUGAG X CGAA ACGUGGGU	1844
1075	CCACGUU C CCAACCUC	70	GAGGUUGG CUGAUGAG X CGAA AACGUGGG	1845
1083	CCAACCU C UUCUCCCU	71	AGGGAGAA CUGAUGAG X CGAA AGGUUGGG	1846
1085	CAACCUCU U CUCCUGC	72	GCAGGGAG CUGAUGAG X CGAA AGAGGUUG	1847
1086	AACCUCUU C UCCUGCA	73	UGCAGGGA CUGAUGAG X CGAA AAGAGGUU	1848
1088	CCUCUUU C CCUGCAGC	74	GCUGCAGG CUGAUGAG X CGAA AGAAGAGG	1849
1098	CUGCAGCU U UGUGGUGC	75	GCACCACA CUGAUGAG X CGAA AGCUGCAG	1850
1099	UGCAGCUU U GUGGUGCU	76	AGCACCAC CUGAUGAG X CGAA AAGCUGCA	1851
1112	UGCUGGCU U CCCCUCU	77	UGAGGGGG CUGAUGAG X CGAA AGCCAGCA	1852
1113	GCUGGCUU C CCCCUCU	78	UUGAGGGG CUGAUGAG X CGAA AAGCCAGC	1853
1119	UUCCCCCU C AACGAGUC	79	GACUGGUU CUGAUGAG X CGAA AGGGGGAA	1854
1127	CAACCAGU C UGAAGUGC	80	GCACUUCA CUGAUGAG X CGAA ACUGGUUG	1855
1142	GCUGGCCU C UGUCGGAG	81	CUCCGACA CUGAUGAG X CGAA AGGCCAGC	1856
1146	GCCUCUGU C GGAGGGAG	82	CUCCCUCC CUGAUGAG X CGAA ACAGAGGC	1857
1161	AGCAUGAU C AUUGGAGG	83	CCUCCAAU CUGAUGAG X CGAA AUCAUGCU	1858
1164	AUGAUCAU U GGAGGUAU	84	AUACCUC CUGAUGAG X CGAA AUGAUCAU	1859
1171	UUGGAGGU A UCGACCAC	85	GUGGUCGA CUGAUGAG X CGAA ACCUCCAA	1860
1173	GGAGGUAU C GACCACUC	86	GAGUGGUC CUGAUGAG X CGAA AUACCUC	1861
1181	CGACCACU C GCUGUACA	87	UGUACAGC CUGAUGAG X CGAA AGUGGUCG	1862
1187	CUCGUGU A CACAGGCA	88	UGCCUGUG CUGAUGAG X CGAA ACAGCGAG	1863
1198	CAGGCAGU C UCUGGUU	89	AUACCAGA CUGAUGAG X CGAA ACUGCCUG	1864
1200	GGCAGUCU C UGUUUAU	90	GUUUAACA CUGAUGAG X CGAA AGACUGCC	1865
1205	UCUCUGGU A UACACCCA	91	UGGUGUUA CUGAUGAG X CGAA ACCAGAGA	1866
1207	UCUGGUU A CACCCAUC	92	GAUGGGUG CUGAUGAG X CGAA AUACCAGA	1867
1215	ACACCAU C CGCGGGGA	93	UCCCGCCG CUGAUGAG X CGAA AUGGGUGU	1868
1229	GGAGUGGU A UUAUGAGG	94	CCUCAUAA CUGAUGAG X CGAA ACCACUCC	1869
1231	AGUGGUU U AUGAGGUG	95	CACCUCU CUGAUGAG X CGAA AUACCACU	1870
1232	GUGGUUU A UGAGGUGA	96	UCACCUCA CUGAUGAG X CGAA AAUACCAC	1871
1242	GAGGUGAU C AUUGUGCG	97	CGCACAUA CUGAUGAG X CGAA AUCACCUC	1872
1245	GUGAUCAU U GUGCGGGU	98	ACCCGCAC CUGAUGAG X CGAA AUGAUCAC	1873

Table 18

1260	GUGGAGAU C AAUGGACA	99	UGUCCAUI CUGAUGAG X CGAA AUCUCCAC	1874
1273	GACAGGAU C UGAAAAUG	100	CAUUUUA CUGAUGAG X CGAA AUCCUGUC	1875
1295	CAAGGAGU A CAACUAUG	101	CAUAGUUG CUGAUGAG X CGAA ACUCCUUG	1876
1301	GUACAACU A UGACAAGA	102	UCUUGUCA CUGAUGAG X CGAA AGUUGUAC	1877
1314	AAGAGCAU U GUGGACAG	103	CUGUCCAC CUGAUGAG X CGAA AUGCUCUU	1878
1338	ACCAACCU U CGUUUGCC	104	GGCAAACG CUGAUGAG X CGAA AGGUUGGU	1879
1339	CCAACCUU C GUUUGCCC	105	GGGCAAAC CUGAUGAG X CGAA AAGGUUGG	1880
1342	ACCUUCGU U UGCCCCAG	106	CUUGGGCA CUGAUGAG X CGAA ACGAAGGU	1881
1343	CCUUCGUU U GCCCAAGA	107	UCUUGGGC CUGAUGAG X CGAA AACGAAGG	1882
1358	GAAAGUGU U UGAAGCUG	108	CAGCUUCA CUGAUGAG X CGAA ACACUUUC	1883
1359	AAAGUGUU U GAAGCUGC	109	GCAGCUUC CUGAUGAG X CGAA AACACUUU	1884
1371	GCUGCAGU C AAAUCCAUI	110	AUGGAUUU CUGAUGAG X CGAA ACUGCAGC	1885
1376	AGUCAAAU C CAUCAAGG	111	CCUUGAUG CUGAUGAG X CGAA AUUUGACU	1886
1380	AAAUCCAUI C AAGGCAGC	112	GCUGCCUU CUGAUGAG X CGAA AUGGAUUU	1887
1391	GGCAGCCU C CUCCACGG	113	CCGUGGAG CUGAUGAG X CGAA AGGCUGCC	1888
1394	AGCCUCCU C CACGGAGA	114	UCUCCGUG CUGAUGAG X CGAA AGGAGGCU	1889
1406	GGAGAAGU U CCCUGAUG	115	CAUCAGGG CUGAUGAG X CGAA ACUUCUCC	1890
1407	GAGAAGUU C CCUGAUGG	116	CCAUCAGG CUGAUGAG X CGAA AACUUCUC	1891
1417	CUGAUGGU U UCUGGCUA	117	UAGCCAGA CUGAUGAG X CGAA ACCAUCAG	1892
1418	UGAUGGUU U CUGGCUAG	118	CUAGCCAG CUGAUGAG X CGAA AACCAUCA	1893
1419	GAUGGUUU C UGGCUAGG	119	CCUAGCCA CUGAUGAG X CGAA AAACCAUC	1894
1425	UUCUGGCU A GGAGAGCA	120	UGCUCUCC CUGAUGAG X CGAA AGCCAGAA	1895
1465	CCACCCCU U GGAACAUI	121	AAUGUUC CUGAUGAG X CGAA AGGGUGG	1896
1473	UGGAACAUI U UCCCCAGU	122	ACUGGGAA CUGAUGAG X CGAA AUGUCCA	1897
1474	GGAACAUI U UCCCAGUC	123	GACUGGGA CUGAUGAG X CGAA AAUGUUC	1898
1475	GAACAUIU U CCCAGUCA	124	UGACUGGG CUGAUGAG X CGAA AAAUGUUC	1899
1476	AACAUIUU C CCAGUCAU	125	AUGACUGG CUGAUGAG X CGAA AAAUGUU	1900
1482	UUCCCAGU C AUCUCACU	126	AGUGAGAU CUGAUGAG X CGAA ACUGGGAA	1901
1485	CCAGUCAU C UCACUCUA	127	UAGAGUGA CUGAUGAG X CGAA AUGACUGG	1902
1487	AGUCAUCU C ACUCUACC	128	GGUAGAGU CUGAUGAG X CGAA AGAUGACU	1903
1491	AUCUCACU C UACCUAUU	129	AUUAGGUA CUGAUGAG X CGAA AGUGAGAU	1904
1493	CUCACUCU A CCUAAUGG	130	CCAUIUAG CUGAUGAG X CGAA AGAGUGAG	1905
1497	CUCUACCU A AUGGGUGA	131	UCACCAUI CUGAUGAG X CGAA AGGUAGAG	1906
1509	GGUGAGGU U ACCAACCA	132	UGGUUGGU CUGAUGAG X CGAA ACCUCACC	1907
1510	GUGAGGUU A CCAACCAG	133	CUGGUUGG CUGAUGAG X CGAA AACUCAC	1908
1520	CAACCAGU C CUUCCGCA	134	UGCGGAAG CUGAUGAG X CGAA ACUGGUUG	1909
1523	CCAGUCCU U CCGCAUCA	135	UGAUGC GG CUGAUGAG X CGAA AGGACUGG	1910
1524	CAGUCCUU C CGCAUCAC	136	GUGAUGC G CUGAUGAG X CGAA AAGACUG	1911
1530	UUCGCAU C ACCAUCCU	137	AGGAUGGU CUGAUGAG X CGAA AUGCGGAA	1912
1536	AUCACCAU C CUUCCGCA	138	UGC GGAAG CUGAUGAG X CGAA AUGGUGAU	1913
1539	ACCAUCCU U CCGCAGCA	139	UGCUGCGG CUGAUGAG X CGAA AGGAUGGU	1914
1540	CCAUCUUU C CGCAGCAA	140	UUGCUGCG CUGAUGAG X CGAA AAGGAUGG	1915
1550	GCAGCAAU A CCUGCGGC	141	GCCG CAGG CUGAUGAG X CGAA AUUGCUGC	1916
1580	GGCCACGU C CCAAGACG	142	CGUCUUGG CUGAUGAG X CGAA ACGUGGCC	1917
1594	ACGACUGU U ACAAGUUU	143	AAACUUGU CUGAUGAG X CGAA ACAGUCGU	1918
1595	CGACUGUU A CAAGUUUG	144	CAAACUUG CUGAUGAG X CGAA AACAGUCG	1919
1601	UUACAAGU U UGCCAUUCU	145	AGAUGGCA CUGAUGAG X CGAA ACUUGUAA	1920
1602	UACAAGUU U GCCAUCUC	146	GAGAUGGC CUGAUGAG X CGAA AACUUGUA	1921
1608	UUUGCAU C UCACAGUC	147	GACUGUGA CUGAUGAG X CGAA AUGGCAAA	1922
1610	UGCCAUCU C ACAGUCAU	148	AUGACUGU CUGAUGAG X CGAA AGAUGGCA	1923
1616	CUCACAGU C AUCCACGG	149	CCGUGGAU CUGAUGAG X CGAA ACUGUGAG	1924

Table 18

1619	ACAGUCAU C CACGGGCA	150	UGCCCCUG CUGAUGAG X CGAA AUGACUGU	1925
1632	GGCACUGU U AUGGGAGC	151	GCUCCCAU CUGAUGAG X CGAA ACAGUGCC	1926
1633	GCACUGUU A UGGGAGCU	152	AGCUCCCA CUGAUGAG X CGAA AACAGUGC	1927
1644	GGAGCUGU U AUCAUGGA	153	UCCAUGAU CUGAUGAG X CGAA ACAGCUCC	1928
1645	GAGCUGUU A UCAUGGAG	154	CUCCAUGA CUGAUGAG X CGAA AACAGCUC	1929
1647	GCUGUUAU C AUGGAGGG	155	CCCUCCAU CUGAUGAG X CGAA AUAACAGC	1930
1658	GGAGGGCU U CUACGUUG	156	CAACGUAG CUGAUGAG X CGAA AGCCCUCC	1931
1659	GAGGGCUU C UACGUUGU	157	ACAACGUA CUGAUGAG X CGAA AAGCCCUC	1932
1661	GGGCUUCU A CGUUGUCU	158	AGACAACG CUGAUGAG X CGAA AGAAGCCC	1933
1665	UUCUACGU U GUCUUGA	159	UCAAGAC CUGAUGAG X CGAA ACGUAGAA	1934
1668	UACGUUGU C UUUGAUCG	160	CGAUCAAA CUGAUGAG X CGAA ACAACGUA	1935
1670	CGUUGUCU U UGAUCGGG	161	CCCGAUCA CUGAUGAG X CGAA AGACAACG	1936
1671	GUUGUCUU U GAUCGGGC	162	GCCCGAUC CUGAUGAG X CGAA AAGACAAC	1937
1675	UCUUUGAU C GGGCCCGA	163	UCGGGCCC CUGAUGAG X CGAA AUCAAAGA	1938
1692	AAACGAAU U GGCUUUGC	164	GCAAAGCC CUGAUGAG X CGAA AUUCGUUU	1939
1697	AAUUGGCU U UGCUGUCA	165	UGACAGCA CUGAUGAG X CGAA AGCCAAUU	1940
1698	AUUGGCUU U GCUGUCAG	166	CUGACAGC CUGAUGAG X CGAA AAGCCAAU	1941
1704	UUUGCUGU C AGCGCUG	167	CAAGCGCU CUGAUGAG X CGAA ACAGCAAA	1942
1711	UCAGCGCU U GCCAUGUG	168	CACAUGGC CUGAUGAG X CGAA AGCGCUGA	1943
1730	CGAUGAGU U CAGGACGG	169	CCGUCCUG CUGAUGAG X CGAA ACUCAUCG	1944
1731	GAUGAGUU C AGGACGGC	170	GCCGUCCU CUGAUGAG X CGAA AACUCAUC	1945
1756	AAGGCCCU U UUGUCACC	171	GGUGACAA CUGAUGAG X CGAA AGGGCCUU	1946
1757	AGGCCCUU U UGUCACCU	172	AGGUGACA CUGAUGAG X CGAA AAGGGCCU	1947
1758	GGCCCUUU U GUCACCUU	173	AAGGUGAC CUGAUGAG X CGAA AAAGGGCC	1948
1761	CCUUUUGU C ACCUUGGA	174	UCCAAGGU CUGAUGAG X CGAA ACAAAAGG	1949
1766	UGUCACCU U GGACAUGG	175	CCAUGUCC CUGAUGAG X CGAA AGGUGACA	1950
1787	CUGUGGCU A CAACAUUC	176	GAAUGUUG CUGAUGAG X CGAA AGCCACAG	1951
1794	UACAACAU U CCACAGAC	177	GUCUGUGG CUGAUGAG X CGAA AUGUUGUA	1952
1795	ACAACAUA C CACAGACA	178	UGUCUGUG CUGAUGAG X CGAA AAUGUUGU	1953
1811	AGAUGAGU C AACCCUCA	179	UGAGGGUU CUGAUGAG X CGAA ACUCAUCU	1954
1818	UCAACCCU C AUGACCAU	180	AUGGUCAU CUGAUGAG X CGAA AGGGUUGA	1955
1827	AUGACCAU A GCCUAUGU	181	ACAUAGGC CUGAUGAG X CGAA AUGGUCAU	1956
1832	CAUAGCCU A UGUCAUGG	182	CCAUGACA CUGAUGAG X CGAA AGGCUAUG	1957
1836	GCCUAUGU C AUGGCUGC	183	GCAGCCAU CUGAUGAG X CGAA ACAUAGGC	1958
1848	GCUGCCAU C UGCGCCCU	184	AGGGCGCA CUGAUGAG X CGAA AUGGCAGC	1959
1857	UGCGCCCU C UUCAUGCU	185	AGCAUGAA CUGAUGAG X CGAA AGGGCGCA	1960
1859	CGCCCUU C AUGCUGC	186	GCAGCAUG CUGAUGAG X CGAA AGAGGGCG	1961
1860	GCCCUUU C AUGCUGCC	187	GGCAGCAU CUGAUGAG X CGAA AAGAGGGC	1962
1872	CUGCCACU C UGCCUCAU	188	AUGAGGCA CUGAUGAG X CGAA AGUGGCAG	1963
1878	CUCUGCCU C AUGGUGUG	189	CACACCAU CUGAUGAG X CGAA AGGCAGAG	1964
1888	UGGUGUGU C AGUGGCGC	190	GCGCCACU CUGAUGAG X CGAA ACACACCA	1965
1902	CGCUGCCU C CGCUGCCU	191	AGGCAGCG CUGAUGAG X CGAA AGGCAGCG	1966
1931	UGAUGACU U UGCUAUG	192	CAUCAGCA CUGAUGAG X CGAA AGUCAUCA	1967
1932	GAUGACUU U GCUGAUGA	193	UCAUCAGC CUGAUGAG X CGAA AAGUCAUC	1968
1944	GAUGACAU C UCCCUGCU	194	AGCAGGGA CUGAUGAG X CGAA AUGUCAUC	1969
1946	UGACAUCU C CCUGCUGA	195	UCAGCAGG CUGAUGAG X CGAA AGAUGUCA	1970
1981	CAGAAGAU A GAGAUUCC	196	GGAAUCUC CUGAUGAG X CGAA AUCUUCUG	1971
1987	AUAGAGAU U CCCUGGA	197	UCCAGGGG CUGAUGAG X CGAA AUCUCUAU	1972
1988	UAGAGAU C CCCUGGAC	198	GUCCAGGG CUGAUGAG X CGAA AAUCUCUA	1973
2004	CCACACCU C CGUGGUUC	199	GAACCACG CUGAUGAG X CGAA AGGUGUGG	1974
2011	UCCGUGGU U CACUUGG	200	CCAAAGUG CUGAUGAG X CGAA ACCACGGA	1975

Table 18

2012	CCGUGGUU C ACUUUGGU	201	ACCAAAGU CUGAUGAG X CGAA AACCACGG	1976
2016	GGUUCACU U UGGUCACA	202	UGUGACCA CUGAUGAG X CGAA AGUGAACC	1977
2017	GUUCACUU U GGUCACAA	203	UUGUGACC CUGAUGAG X CGAA AAGUGAAC	1978
2021	ACUUUGGU C ACAAGUAG	204	CUACUUGU CUGAUGAG X CGAA ACCAAAGU	1979
2028	UCACAAAGU A GGAGACAC	205	GUGUCUCC CUGAUGAG X CGAA ACUUGUGA	1980
2063	GAGCACCU C AGGACCCU	206	AGGGUCCU CUGAUGAG X CGAA AGGUGCUC	1981
2072	AGGACCCU C CCCACCCA	207	UGGGUGGG CUGAUGAG X CGAA AGGGUCCU	1982
2091	AAAUGCCU C UGCCUUGA	208	UCAAGGCA CUGAUGAG X CGAA AGGCAUUU	1983
2097	CUCUGCCU U GAUGGAGA	209	UCUCCAUC CUGAUGAG X CGAA AGGCAGAG	1984
2129	AGGUGGGU U CCAGGGAC	210	GUCCCUUG CUGAUGAG X CGAA ACCCACC	1985
2130	GGUGGGUU C CAGGGACU	211	AGUCCUG CUGAUGAG X CGAA AACCACCC	1986
2141	GGGACUGU A CCUGUAGG	212	CCUACAGG CUGAUGAG X CGAA ACAGUCCC	1987
2147	GUACCUGU A GGAAACAG	213	CUGUUUCC CUGAUGAG X CGAA ACAGGUAC	1988
2177	GAAGCACU C UGCUGGCG	214	CGCCAGCA CUGAUGAG X CGAA AGUGCUC	1989
2191	GCGGGAU A CUCUUGGU	215	ACCAAGAG CUGAUGAG X CGAA AUUCCCGC	1990
2194	GGAAUACU C UUGGUCAC	216	GUGACCAA CUGAUGAG X CGAA AGUAUUC	1991
2196	AAUACUCU U GGUCACCU	217	AGGUGACC CUGAUGAG X CGAA AGAGUAU	1992
2200	CUCUUGGU C ACCUCAA	218	UUUGAGGU CUGAUGAG X CGAA ACCAAGAG	1993
2205	GGUCACCU C AAAUUUAA	219	UUAAAUUU CUGAUGAG X CGAA AGGUGACC	1994
2210	CCUCAAU U UAAGUCGG	220	CCGACUUA CUGAUGAG X CGAA AUUUGAGG	1995
2211	CUCAAAU U AAGUCGGG	221	CCCACUU CUGAUGAG X CGAA AAUUGAG	1996
2212	UCAAAUU A AGUCGGA	222	UCCCGACU CUGAUGAG X CGAA AAAUUUGA	1997
2216	AUUUAAGU C GGAAAUU	223	AAUUUCCC CUGAUGAG X CGAA ACUUAUU	1998
2224	CGGGAUU U CUGCUGCU	224	AGCAGCAG CUGAUGAG X CGAA AUUUCCCC	1999
2225	GGGAUUU C UGCUGCU	225	AAGCAGCA CUGAUGAG X CGAA AAUUUCCC	2000
2233	CUGCUGCU U GAAACUUC	226	GAAGUUUC CUGAUGAG X CGAA AGCAGCAG	2001
2240	UGAAACU U CAGCCUG	227	CAGGGCUG CUGAUGAG X CGAA AGUUUCAA	2002
2241	UGAAACU C AGCCUGA	228	UCAGGGCU CUGAUGAG X CGAA AAGUUUCA	2003
2254	CUGAACCU U UGUCCACC	229	GGUGGACA CUGAUGAG X CGAA AGGUUCAG	2004
2255	UGAACCU U GUCCACCA	230	UGGUGGAC CUGAUGAG X CGAA AAGGUUCA	2005
2258	ACCUUUGU C CACCAUUC	231	GAAUGGUG CUGAUGAG X CGAA ACAAAGGU	2006
2265	UCCACCAU U CCUUUAAA	232	UUUAAAGG CUGAUGAG X CGAA AUGGUGGA	2007
2266	CCACCAU C CUUUAAU	233	AUUUAAAG CUGAUGAG X CGAA AAUGGUGG	2008
2269	CAUUCU U UAAAUUCU	234	AGAAUUUA CUGAUGAG X CGAA AGGAAUGG	2009
2270	CAUUCU U AAAUUCUC	235	GAGAAUUU CUGAUGAG X CGAA AAGGAAUG	2010
2271	AUUCUUU A AAUUCUCC	236	GGAGAAUU CUGAUGAG X CGAA AAAGGAAU	2011
2275	CUUUAUU U CUCCAACC	237	GGUUGGAG CUGAUGAG X CGAA AUUUAAAG	2012
2276	UUUAAAU C UCCAACCC	238	GGGUUGGA CUGAUGAG X CGAA AAUUUAAA	2013
2278	UAAAUUCU C CAACCCAA	239	UUGGGUUG CUGAUGAG X CGAA AGAAUUUA	2014
2290	CCCAAAGU A UUCUUCU	240	AAGAAGAA CUGAUGAG X CGAA ACUUUGGG	2015
2292	CAAAGU U CUUCUUU	241	AAAAGAAG CUGAUGAG X CGAA AUACUUUG	2016
2293	AAAGUAU C UUCUUUUC	242	GAAAAGAA CUGAUGAG X CGAA AAUACUUU	2017
2295	AGUAUUCU U CUUUUCU	243	AAGAAAAG CUGAUGAG X CGAA AGAAUACU	2018
2296	GUUAUCU C UUUUCUUA	244	UAAGAAA CUGAUGAG X CGAA AAGAAUAC	2019
2298	AUUCUUCU U UUCUAGU	245	ACUAAGAA CUGAUGAG X CGAA AGAAGAAU	2020
2299	UUCUUCU U UCUAGUU	246	AACUAAGA CUGAUGAG X CGAA AAGAAGAA	2021
2300	UCUUCUU U CUUAGUU	247	AAACUAG CUGAUGAG X CGAA AAACAAGA	2022
2301	CUUCUUU C UUAGUUUC	248	GAAACUAA CUGAUGAG X CGAA AAAAGAAG	2023
2303	UCUUUUU U AGUUUCAG	249	CUGAAACU CUGAUGAG X CGAA AGAAAAGA	2024
2304	UUUUUCU A GUUCAGA	250	UCUGAAAC CUGAUGAG X CGAA AAGAAAAG	2025
2307	UUCUUAGU U UCAGAAGU	251	ACUUCUGA CUGAUGAG X CGAA ACUAAGAA	2026

Table 18

2308	UCUUGUUU U CAGAAGUA	252	UACUUCUG CUGAUGAG X CGAA AACUAAGA	2027
2309	CUUAGUUU C AGAAGUAC	253	GUACUUCU CUGAUGAG X CGAA AACUAAG	2028
2316	UCAGAAGU A CUGGCAUC	254	GAUGCCAG CUGAUGAG X CGAA ACUUCUGA	2029
2324	ACUGGCAU C ACACGCAG	255	CUGCGUGU CUGAUGAG X CGAA AUGCCAGU	2030
2335	ACGCAGGU U ACCUUGGC	256	GCCAAGGU CUGAUGAG X CGAA ACCUGCGU	2031
2336	CGCAGGUU A CCUUGGCG	257	CGCCAAGG CUGAUGAG X CGAA AACCUGCG	2032
2340	GGUUACCU U GCGUGUG	258	CACACGCC CUGAUGAG X CGAA AGGUAACC	2033
2350	GCGUGUGU C CCUGUGGU	259	ACCACAGG CUGAUGAG X CGAA ACACACGC	2034
2359	CCUGUGGU A CCCUGGCA	260	UGCCAGGG CUGAUGAG X CGAA ACCACAGG	2035
2384	ACCAAGCU U GUUCCCU	261	AGGGAAAC CUGAUGAG X CGAA AGCUUGGU	2036
2387	AAGCUUGU U UCCCUGCU	262	AGCAGGGA CUGAUGAG X CGAA ACAAGCUU	2037
2388	AGCUUGUU U CCCUGCUG	263	CAGCAGGG CUGAUGAG X CGAA AACAAAGCU	2038
2389	GCUUGUUU C CCUGCUG	264	CCAGCAGG CUGAUGAG X CGAA AAACAAGC	2039
2405	GCCAAAGU C AGUAGGAG	265	CUCCUACU CUGAUGAG X CGAA ACUUGGC	2040
2409	AAGUCAGU A GGAGAGGA	266	UCCUCUCC CUGAUGAG X CGAA ACUGACUU	2041
2426	UGCACAGU U UGCUAUUU	267	AAAUAGCA CUGAUGAG X CGAA ACUGUGCA	2042
2427	GCACAGUU U GCUAUUUG	268	CAAAUAGC CUGAUGAG X CGAA AACUGUGC	2043
2431	AGUUGCU A UUUGCUUU	269	AAAGCAAA CUGAUGAG X CGAA AGCAAACU	2044
2433	UUUGCUAU U UGCUUAG	270	CUAAAGCA CUGAUGAG X CGAA AUAGCAAA	2045
2434	UUGCUAUU U GCUUAGA	271	UCUAAAGC CUGAUGAG X CGAA AAUAGCAA	2046
2438	UAUUGCU U UAGAGACA	272	UGUCUCUA CUGAUGAG X CGAA AGCAAUA	2047
2439	AUUUGCUU U AGAGACAG	273	CUGUCUCU CUGAUGAG X CGAA AAGCAAU	2048
2440	UUUGCUUU A GAGACAGG	274	CCUGUCUC CUGAUGAG X CGAA AAAGCAAA	2049
2455	GGGACUGU A UAAACAAG	275	CUUGUUUA CUGAUGAG X CGAA ACAGUCCC	2050
2457	GACUGUUA A AACAAGCC	276	GGCUUGUU CUGAUGAG X CGAA AUACAGUC	2051
2467	ACAAGCCU A ACAUUGGU	277	ACCAAUGU CUGAUGAG X CGAA AGGCUUGU	2052
2472	CCUAACAU U GGUGCAAA	278	UUUGCACC CUGAUGAG X CGAA AUGUUAGG	2053
2484	GCAAAGAU U GCCUCUUG	279	CAAGAGGC CUGAUGAG X CGAA AUCUUUGC	2054
2489	GAUUGCCU C UUGAAUUA	280	UAAUCAA CUGAUGAG X CGAA AGGCAAUC	2055
2491	UUGCCUCU U GAAUUA	281	UUUAAUUC CUGAUGAG X CGAA AGAGGCAA	2056
2496	UCUUGAAU U AAAAAAA	282	UUUUUUUU CUGAUGAG X CGAA AUUCAAGA	2057
2497	CUUGAAUU A AAAAAAA	283	UUUUUUUU CUGAUGAG X CGAA AAUUAAG	2058
2510	AAAAAACU A GAAAAAA	284	UUUUUUUC CUGAUGAG X CGAA AGUUUUUU	2059

Input Sequence = AF190725. Cut Site = G/.

Stem Length = 8. Core Sequence = CUGAUGAG X CGAA (X = GCCGUUAGGC or other stem II)
AF190725 (Homo sapiens beta-site APP cleaving enzyme (BACE) mRNA; 2526 bp)

Table 19

Table 19: Human BACE NCH Ribozyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
10	CACGCGUC C GCAGCCCG	285	CGGGCUGC CUGAUGAG X CGAA IACGCGUG	2060
13	GCGUCCGC A GCCCGCCC	286	GGGCGGGC CUGAUGAG X CGAA ICGGACGC	2061
16	UCCGCAGC C CGCCCGGG	287	CCCGGGCG CUGAUGAG X CGAA ICUGCGGA	2062
17	CCGCAGCC C GCCCGGGA	288	UCCCGGGC CUGAUGAG X CGAA ICGGCGG	2063
20	CAGCCCGC C CGGAGCU	289	AGCUGCCG CUGAUGAG X CGAA ICGGCGUG	2064
21	AGCCCGCC C GGGAGCUG	290	CAGCUGCC CUGAUGAG X CGAA ICGGCGCU	2065
28	CCGGGAGC U GCGAGCCG	291	CGGCUCGC CUGAUGAG X CGAA ICUCCCGG	2066
35	CUGCGAGC C GCGAGCUG	292	CAGCUCGC CUGAUGAG X CGAA ICUCGCAG	2067
42	CCGCGAGC U GGAUUAUG	293	CAUAAUCC CUGAUGAG X CGAA ICUCGCGG	2068
56	AUGGUGGC C UGAGCAGC	294	GCUGCUCA CUGAUGAG X CGAA ICCACCAU	2069
57	UGGUGGCC U GAGCAGCC	295	GGCUGCUC CUGAUGAG X CGAA IGCCACCA	2070
62	GCCUGAGC A GCCAACGC	296	GCGUUGGC CUGAUGAG X CGAA ICUCAGGC	2071
65	UGAGCAGC C AACGCAGC	297	GCUGCGUU CUGAUGAG X CGAA ICUCUCA	2072
66	GAGCAGCC A ACGCAGCC	298	GGCUGCGU CUGAUGAG X CGAA IGCUGCUC	2073
71	GCCAACGC A GCCGAGG	299	CCUGCGGC CUGAUGAG X CGAA ICGUUGGC	2074
74	AACGCAGC C GCAGGAGC	300	GCUCCUGC CUGAUGAG X CGAA ICUGCGUU	2075
77	GCAGCCGC A GGAGCCCG	301	CGGGCUC CUGAUGAG X CGAA ICGGCGUC	2076
83	GCAGGAGC C CGGAGCCC	302	GGGCUCCG CUGAUGAG X CGAA ICUCUCG	2077
84	CAGGAGCC C GGAGCCCU	303	AGGGCUC CUGAUGAG X CGAA IGCUCUG	2078
90	CCCGGAGC C CUUGCCCC	304	GGGGCAAG CUGAUGAG X CGAA ICUCGGG	2079
91	CCGGAGCC C UUGCCCU	305	AGGGCAAG CUGAUGAG X CGAA IGCUCGG	2080
92	CGGAGCCC U UGCCCCUG	306	CAGGGCA CUGAUGAG X CGAA IGCUCGG	2081
96	GCCCUUGC C CCUGCCCG	307	CGGGCAGG CUGAUGAG X CGAA ICAAGGGC	2082
97	CCCUUGCC C CUGCCCGC	308	GCGGGCAG CUGAUGAG X CGAA IGCAAGGG	2083
98	CCUUGCCC C UGCCCGCG	309	CGCGGGCA CUGAUGAG X CGAA IGGCAAGG	2084
99	CUUGCCCC U GCCCGCGC	310	GCGCGGGC CUGAUGAG X CGAA IGGGCAAG	2085
102	GCCCCUGC C CGCGCCGC	311	GCGGCGCG CUGAUGAG X CGAA ICAGGGGC	2086
103	CCCCUGCC C GCGCCGCC	312	GGCGGGCG CUGAUGAG X CGAA IGCAGGGG	2087
108	GCCCGCGC C GCCGCCCG	313	CGGGCGGC CUGAUGAG X CGAA ICGCGGGC	2088
111	CGCGCCGC C GCCGCCCG	314	CGGGCGGC CUGAUGAG X CGAA ICGCGGGC	2089
114	GCCCGCGC C CGCGGGG	315	CCCCGGCG CUGAUGAG X CGAA ICGGCGGC	2090
115	CCGCGGCC C GCCGGGGG	316	CCCCGGCG CUGAUGAG X CGAA ICGGCGGC	2091
118	CCGCGCGC C GGGGGGAC	317	GUCCCCC CUGAUGAG X CGAA ICGGCGGC	2092
127	GGGGGGAC C AGGAAGC	318	GCUUCCCU CUGAUGAG X CGAA IUCCCCC	2093
128	GGGGGACC A GGAAGCC	319	GGCUUCC CUGAUGAG X CGAA IGUCCCC	2094
136	AGGAAGC C GCCACCGG	320	CCGGUGGC CUGAUGAG X CGAA ICUUCCCU	2095
139	GAAGCCGC C ACCGGCCC	321	GGGCCCGU CUGAUGAG X CGAA ICGGCUUC	2096
140	AAGCCGCC A CCGGCCCG	322	CGGGCCGG CUGAUGAG X CGAA ICGGCUU	2097
142	GCCGCCAC C GGCCCGCC	323	GGCGGGCC CUGAUGAG X CGAA IUGGCGGC	2098
146	CCACCGGC C CGCAUGC	324	GCAUGGCG CUGAUGAG X CGAA ICCGGUGG	2099
147	CACCGGCC C GCAUGCC	325	GGCAUGGC CUGAUGAG X CGAA ICGGCGUG	2100
150	CGGCCCGC C AUGCCCGC	326	GCGGGCAU CUGAUGAG X CGAA ICGGCGCG	2101
151	GGCCCGCC A UGCCCGCC	327	GGCGGGCA CUGAUGAG X CGAA ICGGGGCC	2102
155	CGCCAUGC C CGCCCUUC	328	GAGGGGCG CUGAUGAG X CGAA ICAUGGCG	2103
156	GCCAUGCC C GCCCUUC	329	GGAGGGGC CUGAUGAG X CGAA IGCAUGGC	2104
159	AUGCCCGC C CCUCCAG	330	CUGGGAGG CUGAUGAG X CGAA ICGGGCAU	2105
160	UGCCCGCC C CUCCAGC	331	GCUGGGAG CUGAUGAG X CGAA ICGGGCA	2106

Table 19

161	GCCCCCCC C UCCCAGCC	332	GGCUGGGA CUGAUGAG X CGAA IGGCGGGC	2107
162	CCCCGCCC U CCCAGCCC	333	GGGCUGGG CUGAUGAG X CGAA IGGCGGGG	2108
164	CGCCCCUC C CAGCCCCG	334	CGGGGUG CUGAUGAG X CGAA IAGGGGCG	2109
165	GCCCCUCC C AGCCCCGC	335	GCGGGGCU CUGAUGAG X CGAA IGAGGGGC	2110
166	CCCCUCCC A GCCCCGCC	336	GGCGGGGC CUGAUGAG X CGAA IGGAGGGG	2111
169	CUCCCAGC C CCGCCGGG	337	CCCCGGCG CUGAUGAG X CGAA ICUGGGAG	2112
170	UCCCAGCC C CGCCGGGA	338	UCCCGGCG CUGAUGAG X CGAA IGCUGGGA	2113
171	CCCAGCCC C GCCGGGAG	339	CUCCCGGC CUGAUGAG X CGAA IGGCUGGG	2114
174	AGCCCCGC C GGGAGCCC	340	GGGCUCCC CUGAUGAG X CGAA ICGGGGCU	2115
181	CCGGGAGC C CGCCTCCG	341	CGGGGCGG CUGAUGAG X CGAA ICUCCCGG	2116
182	CGGGAGCC C GCGCCCGC	342	GCGGGCGC CUGAUGAG X CGAA IGCUCCCG	2117
187	GCCCCGCG C CGCUGCCC	343	GGGCAGCG CUGAUGAG X CGAA ICGCGGGC	2118
188	CCCGCGCC C GCUGCCCA	344	UGGGCAGC CUGAUGAG X CGAA ICGCGGGG	2119
191	GCGCCCGC U GCCCAGGC	345	GCCUGGGC CUGAUGAG X CGAA ICGGGCGC	2120
194	CCCGCUGC C CAGGCUGG	346	CCAGCCUG CUGAUGAG X CGAA ICAGCGGG	2121
195	CCGCUGCC C AGGCUGGC	347	GCCAGCCU CUGAUGAG X CGAA IGCAGCGG	2122
196	CGCUGCCC A GGCUGGCC	348	GGCCAGCC CUGAUGAG X CGAA IGGCAGCG	2123
200	GCCCAGGC U GGCCGCCG	349	CGGCGGCC CUGAUGAG X CGAA ICCUGGGC	2124
204	AGGCUGGC C GCCCGCGU	350	ACGGCGGC CUGAUGAG X CGAA ICCAGCCU	2125
207	CUGGCCGC C GCCUGGCC	351	GGCACGGC CUGAUGAG X CGAA ICGGCCAG	2126
210	GCCGCCGC C GUGCCGAU	352	AUCGGCAC CUGAUGAG X CGAA ICGGCGGC	2127
215	CGCCGUGC C GAUGUAGC	353	GCUACAUC CUGAUGAG X CGAA ICACGGCG	2128
228	UAGCGGGC U CCGGAUCC	354	GGAUCCGG CUGAUGAG X CGAA ICCCGCUA	2129
230	GCGGGCUC C GGAUCCCA	355	UGGGAUCC CUGAUGAG X CGAA IAGCCCGC	2130
236	UCCGGAUC C CAGCCUCU	356	AGAGGCGU CUGAUGAG X CGAA IAUCCGGA	2131
237	CCGGAUCC C AGCCUCUC	357	GAGAGGCU CUGAUGAG X CGAA IGAUCCGG	2132
238	CGGAUCCC A GCCUCUCC	358	GGAGAGGC CUGAUGAG X CGAA IGGAUCCG	2133
241	AUCCAGCC U UCUCUCCU	359	AGGGGAGA CUGAUGAG X CGAA ICUGGGAU	2134
242	UCCAGGCC U CUCCUCCG	360	CAGGGGAG CUGAUGAG X CGAA IGCUGGGA	2135
244	CCAGCCUC U CCCCUGCU	361	AGCAGGGG CUGAUGAG X CGAA IAGGCUGG	2136
246	AGCCUCUC C CCUGCUCC	362	GGAGCAGG CUGAUGAG X CGAA IAGAGGCU	2137
247	GCCUCUCC C CUGCUCCC	363	GGGAGCAG CUGAUGAG X CGAA IGAGAGGC	2138
248	CCUCUCCC C UGCUCCCG	364	CGGGAGCA CUGAUGAG X CGAA IGGAGAGG	2139
249	CUCUCCCC U GCUCCCGU	365	ACGGGAGC CUGAUGAG X CGAA IGGGAGAG	2140
252	UCCCCUGC U CCCGUGCU	366	AGCACGGG CUGAUGAG X CGAA ICAGGGGA	2141
254	CCUGCUC C CGUGCUCU	367	AGAGCACG CUGAUGAG X CGAA IAGCAGGG	2142
255	CCUGCUCC C GUGCUCUG	368	CAGAGCAC CUGAUGAG X CGAA IGAGCAGG	2143
260	UCCCGUGC U CUGCGGAU	369	AUCCGCAG CUGAUGAG X CGAA ICACGGGA	2144
262	CCGUGCUC U GCGGAUCU	370	AGAUCGCG CUGAUGAG X CGAA IAGCACGG	2145
270	UGCGGAUC U CCCCUGAC	371	GUCAGGGG CUGAUGAG X CGAA IAUCCGCA	2146
272	CGGAUCUC C CCUGACCG	372	CGGUCAGG CUGAUGAG X CGAA IAGAUCCG	2147
273	GGAUCUCC C CUGACCGC	373	GCGGUCAG CUGAUGAG X CGAA IGAGAUCC	2148
274	GAUCUCCC C UGACCGCU	374	AGCGGUCA CUGAUGAG X CGAA IGGAGAUC	2149
275	AUCUCCCC U GACCGCUC	375	GAGCGGUC CUGAUGAG X CGAA IGGGAGAU	2150
279	CCCCUGAC C GCUCUCCA	376	UGGAGAGC CUGAUGAG X CGAA IUCAGGGG	2151
282	CUGACCGC U CUCCACAG	377	CUGUGGAG CUGAUGAG X CGAA ICGGUCAG	2152
284	GACCGCUC U CCACAGCC	378	GGCUGUGG CUGAUGAG X CGAA IAGCGGUC	2153
286	CCGCUCUC C ACAGCCCG	379	CGGGCUGU CUGAUGAG X CGAA IAGAGCGG	2154
287	CGCUCUCC A CAGCCCGG	380	CCGGGUGU CUGAUGAG X CGAA IGAGAGCG	2155
289	CUCUCCAC A GCCCGGAC	381	GUCCGGGC CUGAUGAG X CGAA IUUGAGAG	2156
292	UCCACAGC C CGGACCCG	382	CGGGUCCG CUGAUGAG X CGAA ICUGUGGA	2157

Table 19

293	CCACAGCC C GGACCCGG	383	CCGGGUCC CUGAUGAG X CGAA IGCUGUGG	2158
298	GCCCGGAC C CGGGGCU	384	AGCCCCCG CUGAUGAG X CGAA IUCCGGGC	2159
299	CCCGGACC C GGGGGCUG	385	CAGCCCC CUGAUGAG X CGAA IGUCCGGG	2160
306	CCGGGGGC U GGCCAGG	386	CCUGGGCC CUGAUGAG X CGAA ICCCCCGG	2161
310	GGGCUGGC C CAGGGCCC	387	GGGCCCUG CUGAUGAG X CGAA ICCAGCCC	2162
311	GGCUGGCC C AGGGCCCU	388	AGGGCCCU CUGAUGAG X CGAA IGCCAGCC	2163
312	GCUGGCCC A GGGCCUG	389	CAGGGCCC CUGAUGAG X CGAA IGGCCAGC	2164
317	CCCAGGGC C CUGCAGGC	390	GCCUGCAG CUGAUGAG X CGAA ICCCUGGG	2165
318	CCAGGGCC C UGCAGGCC	391	GGCCUGCA CUGAUGAG X CGAA IGCCUGG	2166
319	CAGGGCCC U GCAGGCC	392	GGGCCUGC CUGAUGAG X CGAA IGGCCCUG	2167
322	GGCCUGC A GGCCUGG	393	CCAGGGCC CUGAUGAG X CGAA ICAGGGCC	2168
326	CUGCAGGC C CUGGCGUC	394	GACGCCAG CUGAUGAG X CGAA ICCUGCAG	2169
327	UGCAGGCC C UGGCGUCC	395	GGACGCCA CUGAUGAG X CGAA IGCCUGCA	2170
328	GCAGGCC U GGCUGCCU	396	AGGACGCC CUGAUGAG X CGAA IGGCCUGC	2171
335	CUGGCGUC C UGAUGCCC	397	GGGCAUCA CUGAUGAG X CGAA IACGCCAG	2172
336	UGGCGUCC U GAUGCCCC	398	GGGGCAUC CUGAUGAG X CGAA IGACGCCA	2173
342	CCUGAUGC C CCAAGCU	399	AGCUUGGG CUGAUGAG X CGAA ICAUCAGG	2174
343	CUGAUGCC C CCAAGCUC	400	GAGCUUGG CUGAUGAG X CGAA IGCAUCAG	2175
344	UGAUGCCC C CAAGCUCC	401	GGAGCUUG CUGAUGAG X CGAA IGGCAUCA	2176
345	GAUGCCCC C AAGCUCCC	402	GGGAGCUU CUGAUGAG X CGAA IGGGCAUC	2177
346	AUGCCCC A AGCUCCCU	403	AGGGAGCU CUGAUGAG X CGAA IGGGCAU	2178
350	CCCCAAGC U CCCUCUCC	404	GGAGAGGG CUGAUGAG X CGAA ICUUGGGG	2179
352	CCAAGCUC C CUCUCCUG	405	CAGGAGAG CUGAUGAG X CGAA IAGCUUGG	2180
353	CAAGCUCC C UCUCUGA	406	UCAGGAGA CUGAUGAG X CGAA IGAGCUUG	2181
354	AAGCUCCC U CUCCUGAG	407	CUCAGGAG CUGAUGAG X CGAA IGGAGCUU	2182
356	GCUCCUC U CCUGAGAA	408	UUCUCAGG CUGAUGAG X CGAA IAGGGAGC	2183
358	UCCUCUC C UGAGAAGC	409	GTUUCUCA CUGAUGAG X CGAA IAGAGGGA	2184
359	CCCUCUC U GAGAAGCC	410	GUUCUCUC CUGAUGAG X CGAA IGAGAGGG	2185
367	UGAGAAGC C ACCAGCAC	411	GUGUCUGU CUGAUGAG X CGAA ICUUCUCA	2186
368	GAGAAGCC A CCAGCACC	412	GGUGCUGG CUGAUGAG X CGAA IGCUCUC	2187
370	GAAGCCAC C AGCACCAC	413	GUGGUGCU CUGAUGAG X CGAA IUGGCUUC	2188
371	AAGCCACC A GCACCACC	414	GGUGGUGC CUGAUGAG X CGAA IGUGGCUU	2189
374	CCACCAGC A CCACCAG	415	CUGGGUGG CUGAUGAG X CGAA ICUGGUGG	2190
376	ACCAGCAC C ACCAGAC	416	GUCUGGGU CUGAUGAG X CGAA IUGCUGGU	2191
377	CCAGCACC A CCCAGACU	417	AGUCUGGG CUGAUGAG X CGAA IGUGCUGG	2192
379	AGCACCAC C CAGACUUG	418	CAAGUCUG CUGAUGAG X CGAA IUGGUGCU	2193
380	GCACCACC C AGACUUGG	419	CCAAGUCU CUGAUGAG X CGAA IGUGGUGC	2194
381	CACCACCC A GACUUGGG	420	CCCAAGUC CUGAUGAG X CGAA IGGUGGUG	2195
385	ACCCAGAC U UGGGGGCA	421	UGCCCCCA CUGAUGAG X CGAA IUCUGGGU	2196
393	UUGGGGGC A GGCGCCAG	422	CUGGCGCC CUGAUGAG X CGAA ICCCCAA	2197
399	GCAGGCGC C AGGGACGG	423	CCGUCCCU CUGAUGAG X CGAA ICGCCUGC	2198
400	CAGGCGCC A GGGACGGA	424	UCCGUCCC CUGAUGAG X CGAA ICGCCUG	2199
416	ACGUGGGC C AGUGCGAG	425	CUCGCACU CUGAUGAG X CGAA ICCACGU	2200
417	CGUGGGCC A GUGCGAGC	426	GCUCGCAC CUGAUGAG X CGAA IGCCACG	2201
426	GUGCGAGC C CAGAGGGC	427	GCCCUCUG CUGAUGAG X CGAA ICUCGCAC	2202
427	UGCAGGCC C AGAGGGCC	428	GGCCUCU CUGAUGAG X CGAA IGUCGCA	2203
428	GCGAGCCC A GAGGGCCC	429	GGGCCCUC CUGAUGAG X CGAA IGGCUCGC	2204
435	CAGAGGGC C CGAAGGCC	430	GGCCUUCG CUGAUGAG X CGAA ICCUCUG	2205
436	AGAGGGCC C GAAGGCCG	431	CGGCCUUC CUGAUGAG X CGAA IGCCUCU	2206
443	CCGAAGGC C GGGGCCCA	432	UGGGCCCC CUGAUGAG X CGAA ICCUUGG	2207
449	GCCGGGGC C CACCAUGG	433	CCAUGGUG CUGAUGAG X CGAA ICCCGGC	2208

Table 19

450	CCGGGGCC C ACCAUGGC	434	GCCAUGGU CUGAUGAG X CGAA IGCCCCGG	2209
451	CGGGGCCC A CCAUGGCC	435	GGCCAUGG CUGAUGAG X CGAA IGGCCCCG	2210
453	GGGCCCAC C AUGGCCCA	436	UGGGCCAU CUGAUGAG X CGAA IUGGGCCC	2211
454	GGCCCAAC A UGGCCCAA	437	UUGGGCCA CUGAUGAG X CGAA IGUGGGCC	2212
459	ACCAUGGC C CAAGCCCU	438	AGGGCUUG CUGAUGAG X CGAA ICCAUGGU	2213
460	CCAUGGCC C AAGCCUG	439	CAGGGCUU CUGAUGAG X CGAA IGCCAUGG	2214
461	CAUGGCCC A AGCCUGC	440	GCAGGGCU CUGAUGAG X CGAA IGGCCAUG	2215
465	GCCCAAGC C CUGCCUG	441	CAGGGCAG CUGAUGAG X CGAA ICUUGGGC	2216
466	CCCAAGCC C UGCCUGG	442	CCAGGGCA CUGAUGAG X CGAA IGCUUGGG	2217
467	CCAAGCCC U GCCCUGG	443	GCCAGGGC CUGAUGAG X CGAA IGGCUUGG	2218
470	AGCCUGC C CUGGCUCC	444	GGAGCCAG CUGAUGAG X CGAA ICAGGGCU	2219
471	GCCUGCC C UGGCUCCU	445	AGGAGCCA CUGAUGAG X CGAA IGCAGGGC	2220
472	CCCUGCCC U GGCUCUG	446	CAGGAGCC CUGAUGAG X CGAA IGGCAGGG	2221
476	GCCUGGC U CCUGUGU	447	ACAGCAGG CUGAUGAG X CGAA ICCAGGGC	2222
478	CCUGGCUC C UGUGUGG	448	CCACAGCA CUGAUGAG X CGAA IAGCCAGG	2223
479	CUGGCUC U GCUGUGA	449	UCCACAGC CUGAUGAG X CGAA IGAGCCAG	2224
482	GCUCUGC U GUGGAUGG	450	CCAUCCAC CUGAUGAG X CGAA ICAGGAGC	2225
503	GGGAGUGC U GCCUGCC	451	GGGCAGGC CUGAUGAG X CGAA ICACUCCC	2226
506	AGUCUGC U UGCCACG	452	CGUGGGCA CUGAUGAG X CGAA ICAGCACU	2227
507	GUGCUGC U GCCACGG	453	CCGUGGGC CUGAUGAG X CGAA IGCGACAC	2228
510	UGCCUGC C CACGAC	454	GUGCCGUG CUGAUGAG X CGAA ICAGGCAG	2229
511	UGCCUGC C ACGGCACC	455	GGUGCCGU CUGAUGAG X CGAA IGCGAGCA	2230
512	GCCUGCCC A CGGCACCC	456	GGGUGCCG CUGAUGAG X CGAA IGGCAGGC	2231
517	CCCACGGC A CCCAGCAC	457	GUGCUGGG CUGAUGAG X CGAA ICCGUGGG	2232
519	CACGCAC C CAGCACGG	458	CCGUGCUG CUGAUGAG X CGAA IUGCCGUG	2233
520	ACGGACCC C AGCACGGC	459	GCCGUGCU CUGAUGAG X CGAA IGUGCCGU	2234
521	CGGCACCC A GCACGGCA	460	UGCCGUGC CUGAUGAG X CGAA IGGUGCCG	2235
524	CACCCAGC A CGGCAUCC	461	GGAUGCCG CUGAUGAG X CGAA ICUGGGUG	2236
529	AGCACGGC A UCCGGCUG	462	CAGCCGGA CUGAUGAG X CGAA ICCGUGCU	2237
532	ACGGCAUC C GGCUGCC	463	GGGCAGCC CUGAUGAG X CGAA IAUGCCGU	2238
536	CAUCCGGC U GCCCUGC	464	GCAGGGGC CUGAUGAG X CGAA ICCGGAUG	2239
539	CCGGCUGC C CCUGCGA	465	UGCACAGG CUGAUGAG X CGAA ICAGCCGG	2240
540	CGGCUGC C CUGCGCAG	466	CUGCGCAG CUGAUGAG X CGAA IGCGCCG	2241
541	GGCUGCCC C UGCGCAGC	467	GCUGCGCA CUGAUGAG X CGAA IGGCAGCC	2242
542	GCUGCCC U GCGCAGC	468	CGCUGCGC CUGAUGAG X CGAA IGGGCAGC	2243
547	CCCUGCGC A GCGGCCUG	469	CAGGCCGC CUGAUGAG X CGAA ICGCAGGG	2244
553	GCAGCGGC C UGGGGGGC	470	GCCCCCA CUGAUGAG X CGAA ICCGUGC	2245
554	CAGCGGC U GGGGGGCG	471	CGCCCCC CUGAUGAG X CGAA IGCCGUG	2246
564	GGGGGCGC C CCCUGGG	472	CCCAGGGG CUGAUGAG X CGAA ICGCCCC	2247
565	GGGGCGCC C CCCUGGG	473	CCCAGGG CUGAUGAG X CGAA ICGCCCC	2248
566	GGGGCGCC C CCUGGGG	474	GCCCCAGG CUGAUGAG X CGAA IGGCGCC	2249
567	GGCGCCCC C CUGGGGCU	475	AGCCCCAG CUGAUGAG X CGAA IGGCGCC	2250
568	GCGCCCCC C UGGGGCUG	476	CAGCCCCA CUGAUGAG X CGAA IGGGGCG	2251
569	CGCCCCC U GGGGCUG	477	GCAGCCCC CUGAUGAG X CGAA IGGGGCG	2252
575	CCUGGGG U GCGGCUG	478	GCAGCCG CUGAUGAG X CGAA ICCGAGG	2253
581	GCUGCGG U GCCCCGG	479	CCCGGGG CUGAUGAG X CGAA ICCGAGC	2254
584	GCGGCUGC C CCGGAGA	480	UCUCCGG CUGAUGAG X CGAA ICAGCCG	2255
585	CGGCUGC C CCGGAGAC	481	GUCUCCG CUGAUGAG X CGAA IGCAGCC	2256
586	GGCUGCCC C GGGAGACC	482	GGUCUCC CUGAUGAG X CGAA IGGCAGC	2257
594	CGGAGAC C GACGAAGA	483	UCUUCGUC CUGAUGAG X CGAA IUCUCCG	2258
605	CGAAGAGC C CGAGGAGC	484	GCUCCUG CUGAUGAG X CGAA ICUCUCC	2259

Table 19

606	GAAGAGCC C GAGGAGCC	485	GGCUCCUC CUGAUGAG X CGAA IGCUCUUC	2260
614	CGAGGAGC C CGGCCGGA	486	UCCGGCCG CUGAUGAG X CGAA ICUCCUCG	2261
615	GAGGAGCC C GGCCGGAG	487	CUCCGGCC CUGAUGAG X CGAA IGCUCUUC	2262
619	AGCCCGGC C GGAGGGGC	488	GCCCCUCC CUGAUGAG X CGAA ICCGGGCU	2263
628	GGAGGGGC A GCUUGUG	489	CACAAAGC CUGAUGAG X CGAA ICCCUCC	2264
631	GGGGCAGC U UUGUGGAG	490	CUCCACAA CUGAUGAG X CGAA ICUGCCCC	2265
649	UGGUGGAC A ACCUGAGG	491	CCUCAGGU CUGAUGAG X CGAA IUCCACCA	2266
652	UGGACAAC C UGAGGGGC	492	GCCCCUCA CUGAUGAG X CGAA IUUGUCCA	2267
653	GGACAACC U GAGGGGCA	493	UGCCCCUC CUGAUGAG X CGAA IGUUGUCC	2268
661	UGAGGGGC A AGUCGGGG	494	CCCCGACU CUGAUGAG X CGAA ICCCUCA	2269
671	GUCGGGGC A GGGCUACU	495	AGUAGCCC CUGAUGAG X CGAA ICCCGAC	2270
676	GGCAGGGC U ACUACGUG	496	CACGUAGU CUGAUGAG X CGAA ICCUGCC	2271
679	AGGGCUAC U ACGUGGAG	497	CUCCACGU CUGAUGAG X CGAA IUAGCCCU	2272
693	GAGAUGAC C GUGGGCAG	498	CUGCCAC CUGAUGAG X CGAA IUCAUCUC	2273
700	CCGUGGGC A GCCCCCG	499	CGGGGGGC CUGAUGAG X CGAA ICCACGG	2274
703	UGGGCAGC C CCGCGCAG	500	CUGCGGGG CUGAUGAG X CGAA ICUGCCCA	2275
704	GGGAGGCC C CCGCAGA	501	UCUGCGGG CUGAUGAG X CGAA IGCUGCCC	2276
705	GGCAGCCC C CCGCAGAC	502	GUCUGCGG CUGAUGAG X CGAA IGGCUGCC	2277
706	GCAGCCCC C CGCAGACG	503	CGUCUGCG CUGAUGAG X CGAA IGGGUGGC	2278
707	CAGCCCC C GCAGACGC	504	GCGUCUGC CUGAUGAG X CGAA IGGGGCUG	2279
710	CCCCCGC A GACGCUCA	505	UGAGCGUC CUGAUGAG X CGAA ICGGGGGG	2280
716	GCAGACGC U CAACAUCC	506	GGAUGUUG CUGAUGAG X CGAA ICGUCUGC	2281
718	AGACGCUC A ACAUCCUG	507	CAGGAUGU CUGAUGAG X CGAA IAGCGUCU	2282
721	CGCUCAAC A UCCUGGUG	508	CACCAGGA CUGAUGAG X CGAA IUUGAGCG	2283
724	UCAACAUC C UGGUGGAU	509	AUCCACCA CUGAUGAG X CGAA IAUUGUA	2284
725	CAACAUCC U GGUGGAUA	510	UAUCCACC CUGAUGAG X CGAA IGAUGUUG	2285
735	GUGGAUAC A GGCAGCAG	511	CUGCUGCC CUGAUGAG X CGAA IUAUCCAC	2286
739	AUACAGGC A GCAGUAAC	512	GUUACUGC CUGAUGAG X CGAA ICCUGUAU	2287
742	CAGGCAGC A GUAACUUU	513	AAAGUUAC CUGAUGAG X CGAA ICUGCCUG	2288
748	GCAGUAAC U UUGCAGUG	514	CACUGCAA CUGAUGAG X CGAA IUUACUGC	2289
753	AACUUUGC A GUGGGUGC	515	GCACCCAC CUGAUGAG X CGAA ICAAAGUU	2290
762	GUGGGUGC U GCCCCCA	516	UGGGGGGC CUGAUGAG X CGAA ICACCCAC	2291
765	GGUGCUGC C CCCACCC	517	GGGUGGGG CUGAUGAG X CGAA ICAGCACC	2292
766	GUGCUGCC C CCCACCC	518	GGGGUGGG CUGAUGAG X CGAA IGCAGCAC	2293
767	UGCUGCCC C CCACCCU	519	AGGGGUGG CUGAUGAG X CGAA IGGCAGCA	2294
768	GCUGCCCC C CACCCU	520	AAGGGGUG CUGAUGAG X CGAA IGGGCAGC	2295
769	CUGCCCC C ACCCUUC	521	GAAGGGGU CUGAUGAG X CGAA IGGGCAG	2296
770	UGCCCCC A CCCCUC	522	GGAAGGGG CUGAUGAG X CGAA IGGGGCA	2297
772	CCCCCAC C CCUCCUG	523	CAGGAAGG CUGAUGAG X CGAA IUGGGGG	2298
773	CCCCACC C CUUCCUGC	524	GCAGGAAG CUGAUGAG X CGAA IGUGGGG	2299
774	CCCCACC C UCCUGCA	525	UGCAGGAA CUGAUGAG X CGAA IGGUGGG	2300
775	CCCACCC U UCCUGAU	526	AUGCAGGA CUGAUGAG X CGAA IGGUGGG	2301
778	ACCCUUC C UGCAUCGC	527	GCGAUGCA CUGAUGAG X CGAA IAAGGGGU	2302
779	CCCUUCC U GCAUCGCU	528	AGCGAUGC CUGAUGAG X CGAA IGAAGGG	2303
782	CUUCCUGC A UCGCUACU	529	AGUAGCGA CUGAUGAG X CGAA ICAGGAAG	2304
787	UGCAUCGC U ACUACCAG	530	CUGGUAGU CUGAUGAG X CGAA ICAUGCA	2305
790	AUCGCUAC U ACCAGAGG	531	CCUCUGGU CUGAUGAG X CGAA IUAGCGAU	2306
793	GCUACUAC C AGAGGCAG	532	CUGCCUCU CUGAUGAG X CGAA IUAGUAGC	2307
794	CUACUACC A GAGGCAGC	533	GCUGCCUC CUGAUGAG X CGAA IGUAGUAG	2308
800	CCAGAGGC A GCUGUCCA	534	UGGACAGC CUGAUGAG X CGAA ICCUCUGG	2309
803	GAGGCAGC U GUCCAGCA	535	UGCUGGAC CUGAUGAG X CGAA ICUGCCUC	2310

Table 19

807	CAGCUGUC C AGCACAUA	536	UAUGUGCU CUGAUGAG X CGAA IACAGCUG	2311
808	AGCUGUCC A GCACAUAC	537	GUAUGUGC CUGAUGAG X CGAA IGACAGCU	2312
811	UGUCCAGC A CAUACCGG	538	CCGGUAUG CUGAUGAG X CGAA ICUGGACA	2313
813	UCCAGCAC A UACCGGGA	539	UCCCGGUA CUGAUGAG X CGAA IUGCUGGA	2314
817	GCACAUAC C GGGACCUC	540	GAGGUCCC CUGAUGAG X CGAA IUAUGUGC	2315
823	ACCGGGAC C UCCGGAAG	541	CUUCCGGA CUGAUGAG X CGAA IUCCCGGU	2316
824	CCGGGACC U CCGGAAGG	542	CCUCCGG CUGAUGAG X CGAA IGUCCCGG	2317
826	GGGACCUC C GGAAGGGU	543	ACCCUCC CUGAUGAG X CGAA IAGGUCCC	2318
845	GUAUGUGC C CUACACCC	544	GGGUGUAG CUGAUGAG X CGAA ICACAUAC	2319
846	UAUGUGCC C UACACCCA	545	UGGGUGUA CUGAUGAG X CGAA IGCACAUA	2320
847	AUGUGCCC U ACACCCAG	546	CUGGGUGU CUGAUGAG X CGAA IGGCACAU	2321
850	UGCCCUAC A CCCAGGGC	547	GCCUGGG CUGAUGAG X CGAA IUAGGGCA	2322
852	CCCUACAC C CAGGGCAA	548	UUGCCUG CUGAUGAG X CGAA IUGUAGGG	2323
853	CCUACACC C AGGGCAAG	549	CUUGCCCU CUGAUGAG X CGAA IGUGUAGG	2324
854	CUACACCC A GGGCAAGU	550	ACUUGCCC CUGAUGAG X CGAA IGGUGUAG	2325
859	CCCAGGGC A AGUGGGAA	551	UUCCACU CUGAUGAG X CGAA ICCUGGG	2326
875	AGGGGAGC U GGGCACCG	552	CGUGGCC CUGAUGAG X CGAA ICUCCCU	2327
880	AGCUGGGC A CCGACCUG	553	CAGGUCGG CUGAUGAG X CGAA ICCAGCU	2328
882	CUGGGCAC C GACCUGGU	554	ACCAGGUC CUGAUGAG X CGAA IUGCCAG	2329
886	GCACCGAC C UGUAAAGC	555	GCUUACCA CUGAUGAG X CGAA IUCGGUC	2330
887	CACCGACC U GGUAAAGCA	556	UGCUUACC CUGAUGAG X CGAA IGUCGGUG	2331
895	UGGUAAAGC A UCCCCAU	557	AUGGGGGA CUGAUGAG X CGAA ICUUACCA	2332
898	UAAGCAUC C CCCAUGGC	558	GCCAUGGG CUGAUGAG X CGAA IAUUCUUA	2333
899	AAGCAUCC C CCAUGGCC	559	GGCCAUGG CUGAUGAG X CGAA IGAUCUU	2334
900	AGCAUCCC C CAUGGCCC	560	GGGCAUG CUGAUGAG X CGAA IGAUGCU	2335
901	GCAUCCCC C AUGGCCCC	561	GGGGCCAU CUGAUGAG X CGAA IGGGAUGC	2336
902	CAUCCCC A UGGCCCCA	562	UGGGGCCA CUGAUGAG X CGAA IGGGAUG	2337
907	CCCAUGGC C CCAACGUC	563	GACGUUG CUGAUGAG X CGAA ICCAUGGG	2338
908	CCAUGGCC C CAACGUCA	564	UGACGUUG CUGAUGAG X CGAA IGCCAUGG	2339
909	CAUGGCCC C AACGUCAC	565	GUGACGUU CUGAUGAG X CGAA IGGCCAUG	2340
910	AUGGCCCC A ACGUCACU	566	AGUGACGU CUGAUGAG X CGAA IGGCCAUG	2341
916	CCAACGUC A CUGUGCGU	567	ACGCACAG CUGAUGAG X CGAA IACGUUGG	2342
918	AACGUCAC U GUGCGUGC	568	GCACGCAC CUGAUGAG X CGAA IUGACGUU	2343
927	GUGCGUGC C AACAUUGC	569	GCAAUGUU CUGAUGAG X CGAA ICACGCAC	2344
928	UGCUGGCC A ACAUUGCU	570	AGCAAUGU CUGAUGAG X CGAA IGCACGCA	2345
931	GUGCCAAC A UUGCUGCC	571	GGCAGCAA CUGAUGAG X CGAA IUUGGCAC	2346
936	AACAUUGC U GCCAUCAC	572	GUGAUGGC CUGAUGAG X CGAA ICAAUGUU	2347
939	AUUGCUGC C AUCACUGA	573	UCAGUGAU CUGAUGAG X CGAA ICAGCAAU	2348
940	UUGCUGCC A UCACUGAA	574	UUCAGUGA CUGAUGAG X CGAA IGCAGCAA	2349
943	CUGCCAUC A CUGAAUCA	575	UGAUUCAG CUGAUGAG X CGAA IAUGGCAG	2350
945	GCCAUCAC U GAAUCAGA	576	UCUGAUUC CUGAUGAG X CGAA IUGAUGGC	2351
951	ACUGAAUC A GACAAGUU	577	AACUUGUC CUGAUGAG X CGAA IAUUCAGU	2352
955	AAUCAGAC A AGUUCUUC	578	GAAGAACU CUGAUGAG X CGAA IUCUGAUU	2353
961	ACAAGUUC U UCAUCAAC	579	GUUGAUGA CUGAUGAG X CGAA IAACUUGU	2354
964	AGUUCUUC A UCAACGGC	580	GCCGUUGA CUGAUGAG X CGAA IAAGAACU	2355
967	UCUUCAUC A ACGGCUCC	581	GGAGCCGU CUGAUGAG X CGAA IAUGAAGA	2356
973	UCAACGGC U CCAACUGG	582	CCAGUUGG CUGAUGAG X CGAA ICCGUUGA	2357
975	AACGGCUC C AACUGGGA	583	UCCAGUU CUGAUGAG X CGAA IAGCCGUU	2358
976	ACGGCUCC A ACUGGGAA	584	UUCCAGU CUGAUGAG X CGAA IGAGCCGU	2359
979	GCUCCAAC U GGAAGGGC	585	GCCUCCCC CUGAUGAG X CGAA IUUGGAGC	2360
988	GGGAAGGC A UCCUGGGG	586	CCCCAGGA CUGAUGAG X CGAA ICCUCCCC	2361

Table 19

991	AAGGCAUC C UGGGGCUG	587	CAGCCCCA CUGAUGAG X CGAA IAUGCCUU	2362
992	AGGCAUCC U GGGGCUGG	588	CCAGCCCC CUGAUGAG X CGAA IGAUGCCU	2363
998	CCUGGGGC U GGCCUAUG	589	CAUAGGCC CUGAUGAG X CGAA ICCCCAGG	2364
1002	GGGCUGGC C UAUGCUGA	590	UCAGCAUA CUGAUGAG X CGAA ICCAGCCC	2365
1003	GGCUGGCC U AUGCUGAG	591	CUCAGCAU CUGAUGAG X CGAA IGCCAGCC	2366
1008	GCCUAUGC U GAGAUUGC	592	GCAAUCUC CUGAUGAG X CGAA ICAUAGGC	2367
1017	GAGAUUGC C AGGCCUGA	593	UCAGGCCU CUGAUGAG X CGAA ICAAUCUC	2368
1018	AGAUUGCC A GGCCUGAC	594	GUCAGGCC CUGAUGAG X CGAA IGCAAUCU	2369
1022	UGCCAGGC C UGACGACU	595	AGUCGUCA CUGAUGAG X CGAA ICCUGGCA	2370
1023	GCCAGGCC U GACGACUC	596	GAGUCGUC CUGAUGAG X CGAA IGCCUGGC	2371
1030	CUGACGAC U CCCUGGAG	597	CUCCAGGG CUGAUGAG X CGAA IUCGUCAG	2372
1032	GACGACUC C CUGGAGCC	598	GGCUCCAG CUGAUGAG X CGAA IAGUCGUC	2373
1033	ACGACUCC C UGGAGCCU	599	AGGCUCCA CUGAUGAG X CGAA IGAGUCGU	2374
1034	CGACUCCC U GGAGCCUU	600	AAGGCUCC CUGAUGAG X CGAA IGGAGUCG	2375
1040	CCUGGAGC C UUUCUUUG	601	CAAAGAAA CUGAUGAG X CGAA ICUCCAGG	2376
1041	CUGGAGCC U UUCUUUGA	602	UCAAAGAA CUGAUGAG X CGAA IGCUCAG	2377
1045	AGCCUUUC U UUGACUCU	603	AGAGUCAA CUGAUGAG X CGAA IAAAGGCU	2378
1051	UCUUUGAC U CUCUGGUA	604	UACCAGAG CUGAUGAG X CGAA IUCAAAGA	2379
1053	UUUGACUC U CUGGUAAA	605	UUUACCAG CUGAUGAG X CGAA IAGUCAA	2380
1055	UGACUCUC U GGUAAAGC	606	GCUUUACC CUGAUGAG X CGAA IAGAGUCA	2381
1064	GGUAAAGC A GACCCACG	607	CGUGGGUC CUGAUGAG X CGAA ICUUUACC	2382
1068	AAGCAGAC C CACGUUCC	608	GGAACGUG CUGAUGAG X CGAA IUCUGCUU	2383
1069	AGCAGACC C ACGUCC	609	GGGAACGU CUGAUGAG X CGAA IGUCUGCU	2384
1070	GCAGACCC A CGUCCCA	610	UGGGAACG CUGAUGAG X CGAA IGGUCUGC	2385
1076	CCACGUUC C CAACCUCU	611	AGAGGUUG CUGAUGAG X CGAA IAACGUGG	2386
1077	ACGUUCC C AACCUCU	612	AAGAGGUU CUGAUGAG X CGAA IGAACGUG	2387
1078	ACGUUCCC A ACCUCUUC	613	GAAGAGGU CUGAUGAG X CGAA IGGAACGU	2388
1081	UUCCCAAC C UCUCUCC	614	GGAGAAGA CUGAUGAG X CGAA IUUGGGAA	2389
1082	UCCCAACC U CUUCUCC	615	GGGAGAAG CUGAUGAG X CGAA IGUUGGGA	2390
1084	CCAACCUC U UCUCUCC	616	CAGGGAGA CUGAUGAG X CGAA IAGGUUGG	2391
1087	ACCUCUUC U CCCUGCAG	617	CUGCAGGG CUGAUGAG X CGAA IAAGAGGU	2392
1089	CUCUUCUC C CUGCAGCU	618	AGCUGCAG CUGAUGAG X CGAA IAGAAGAG	2393
1090	UCUUCUCC C UGCAGCUU	619	AAGCUGCA CUGAUGAG X CGAA IGAGAAGA	2394
1091	CUUCUCCC U GCAGCUU	620	AAAGCUGC CUGAUGAG X CGAA IGGAGAAG	2395
1094	CUCCUUGC A GCUUUGUG	621	CACAAAGC CUGAUGAG X CGAA ICAGGGAG	2396
1097	CCUGCAGC U UUGUGGUG	622	CACCACAA CUGAUGAG X CGAA ICUGCAGG	2397
1107	UGUGGUGC U GGUUCCC	623	GGGAAGCC CUGAUGAG X CGAA ICACCACA	2398
1111	GUGCUGGC U UCCCCUC	624	GAGGGGGA CUGAUGAG X CGAA ICCAGCAC	2399
1114	CUGGCUUC C CCUCAAC	625	GUUGAGGG CUGAUGAG X CGAA IAAGCCAG	2400
1115	UGGCUUCC C CCUCAACC	626	GGUUGAGG CUGAUGAG X CGAA IGAAGCCA	2401
1116	GGCUUCCC C CUCAACCA	627	UGGUUGAG CUGAUGAG X CGAA IGGAAGCC	2402
1117	GUUCCCC C UCAACCAG	628	CUGGUUGA CUGAUGAG X CGAA IGGGAAGC	2403
1118	CUUCCCC U CAACCAGU	629	ACUGGUUG CUGAUGAG X CGAA IGGGGAAG	2404
1120	UCCCCUC A ACCAGUCU	630	AGACUGGU CUGAUGAG X CGAA IAGGGGGA	2405
1123	CCCUCAAC C AGUCUGAA	631	UUCAGACU CUGAUGAG X CGAA IUUGAGGG	2406
1124	CCUCAACC A GUCUGAAG	632	CUUCAGAC CUGAUGAG X CGAA IGUUGAGG	2407
1128	AACCAGUC U GAAGUGCU	633	AGCACUUC CUGAUGAG X CGAA IACUGGUU	2408
1136	UGAAGUGC U GGCCUCUG	634	CAGAGGCC CUGAUGAG X CGAA ICACUUCA	2409
1140	GUGCUGGC C UCUGUCGG	635	CCGACAGA CUGAUGAG X CGAA ICCAGCAC	2410
1141	UGCUGGCC U CUGUCGGA	636	UCCGACAG CUGAUGAG X CGAA IGCCAGCA	2411
1143	CUGGCCUC U GUCGGAGG	637	CCUCCGAC CUGAUGAG X CGAA IAGGCCAG	2412

Table 19

1156	GAGGGAGC A UGAUCAUU	638	AAUGAUCA CUGAUGAG X CGAA ICUCCUC	2413
1162	GCAUGAUC A UUGGAGGU	639	ACCUCCAA CUGAUGAG X CGAA IAUCAUGC	2414
1177	GUUUCGAC C ACUCGUG	640	CAGCGAGU CUGAUGAG X CGAA IUCGAUAC	2415
1178	UAUCGACC A CUCGUGU	641	ACAGCGAG CUGAUGAG X CGAA IGUCGAUA	2416
1180	UCGACCAC U CGCUGUAC	642	GUACAGCG CUGAUGAG X CGAA IUGGUCGA	2417
1184	CCACUCGC U GUACACAG	643	CUGUGUAC CUGAUGAG X CGAA ICGAGUGG	2418
1189	CGCUGUAC A CAGGCAGU	644	ACUGCCUG CUGAUGAG X CGAA IUACAGCG	2419
1191	CUGUACAC A GGCAGUCU	645	AGACUGCC CUGAUGAG X CGAA IUGUACAG	2420
1195	ACACAGGC A GUCUCUGG	646	CCAGAGAC CUGAUGAG X CGAA ICCUGUGU	2421
1199	AGGCAGUC U CUGGUAUA	647	UAUACCAG CUGAUGAG X CGAA IACUGCCU	2422
1201	GCAGUCUC U GGUUAUACA	648	UGUAUACC CUGAUGAG X CGAA IAGACUGC	2423
1209	UGGUAUAC A CCAUCCG	649	CGGAUGGG CUGAUGAG X CGAA IUUAUACCA	2424
1211	GUUAUACAC C CAUCCGGC	650	GCCGGAUG CUGAUGAG X CGAA IUGUAUAC	2425
1212	UAUACACC C AUCCGGCG	651	CGCCGGAU CUGAUGAG X CGAA IGUGUAUA	2426
1213	AUACACCC A UCCGGCGG	652	CCGCCGGA CUGAUGAG X CGAA IGGUGUAU	2427
1216	CACCAUC C GCGGGAG	653	CUCCCGCC CUGAUGAG X CGAA IAUGGGUG	2428
1243	AGGUGAUC A UUGUGCGG	654	CCGCACAA CUGAUGAG X CGAA IAUCACCU	2429
1261	UGGAGAUC A AUGGACAG	655	CUGUCCAU CUGAUGAG X CGAA IAUCUCCA	2430
1268	CAAUGGAC A GGAUCUGA	656	UCAGAUC CUGAUGAG X CGAA IUCCAUG	2431
1274	ACAGGAUC U GAAAUGG	657	CCAUUUUC CUGAUGAG X CGAA IAUCUGU	2432
1285	AAAUGGAC U GCAAGGAG	658	CUCCUUGC CUGAUGAG X CGAA IUCCAUUU	2433
1288	UGGACUGC A AGGAGUAC	659	GUACUCCU CUGAUGAG X CGAA ICAGUCCA	2434
1297	AGGAGUAC A ACUAUGAC	660	GUCAUAGU CUGAUGAG X CGAA IUACUCCU	2435
1300	AGUACAAC U AUGACAAG	661	CUUGUCAU CUGAUGAG X CGAA IUUGUACU	2436
1306	ACUAUGAC A AGAGCAUU	662	AAUGUCU CUGAUGAG X CGAA IUCAUAGU	2437
1312	ACAAGAGC A UUGUGGAC	663	GUCCACAA CUGAUGAG X CGAA ICUCUUGU	2438
1321	UUGUGGAC A GUGGCACC	664	GGUGCCAC CUGAUGAG X CGAA IUCCACAA	2439
1327	ACAGUGGC A CCACCAAC	665	GUUGGUGG CUGAUGAG X CGAA ICCACUGU	2440
1329	AGUGGCAC C ACCAACCU	666	AGGUUGGU CUGAUGAG X CGAA IUGCCACU	2441
1330	GUGGCACC A CCAACCUU	667	AAGGUUGG CUGAUGAG X CGAA IGUGCCAC	2442
1332	GGCACCAC C AACCUUCG	668	CGAAGGUU CUGAUGAG X CGAA IUGGUGCC	2443
1333	GCACCACC A ACCUUCGU	669	ACGAAGGU CUGAUGAG X CGAA IGUGGUGC	2444
1336	CCACCAAC C UUCGUUUG	670	CAAACGAA CUGAUGAG X CGAA IUUGGUGG	2445
1337	CACCAACC U UCGUUUGC	671	GCAAACGA CUGAUGAG X CGAA IGUUGGUG	2446
1346	UCGUUUGC C CAAGAAAG	672	CUUUCUUG CUGAUGAG X CGAA ICAAACGA	2447
1347	CGUUUGCC C AAGAAAGU	673	ACUUUCUU CUGAUGAG X CGAA IGCAAACG	2448
1348	GUUUUGCC A AGAAAGUG	674	CACUUUCU CUGAUGAG X CGAA IGGCAAAC	2449
1365	UUUGAAGC U GCAGUCAA	675	UUGACUGC CUGAUGAG X CGAA ICUUCAA	2450
1368	GAAGCUGC A GUCAAAUC	676	GAUUUGAC CUGAUGAG X CGAA ICAGCUUC	2451
1372	CUGCAGUC A AAUCCAUC	677	GAUGGAUU CUGAUGAG X CGAA IACUGCAG	2452
1377	GUCAAAUC C AUCAAGGC	678	GCCUUGAU CUGAUGAG X CGAA IAUUUGAC	2453
1378	UCAAAUCC A UCAAGGCA	679	UGCCUUGA CUGAUGAG X CGAA IGAUUUGA	2454
1381	AAUCCAUC A AGGCAGCC	680	GGCUGCCU CUGAUGAG X CGAA IAUGGAUU	2455
1386	AUCAAGGC A GCCUCCUC	681	GAGGAGGC CUGAUGAG X CGAA ICCUUGAU	2456
1389	AAGGCAGC C UCCUCCAC	682	GUGGAGGA CUGAUGAG X CGAA ICUGCCUU	2457
1390	AGGCAGCC U CCUCCACG	683	CGUGGAGG CUGAUGAG X CGAA IGCUGCCU	2458
1392	GCAGCCUC C UCCACGGA	684	UCCGUGGA CUGAUGAG X CGAA IAGGCUGC	2459
1393	CAGCCUCC U CCACGGAG	685	CUCCGUGG CUGAUGAG X CGAA IGAGGCUG	2460
1395	GCCUCCUC C ACGAGAGAA	686	UUCUCCGU CUGAUGAG X CGAA IAGGAGGC	2461
1396	CCUCCUCC A CGGAGAAG	687	CUUCUCCG CUGAUGAG X CGAA IGAGGAGG	2462
1408	AGAAGUUC C CUGAUGGU	688	ACCAUCAG CUGAUGAG X CGAA IAACUUCU	2463

Table 19

1409	GAAGUUC C UGAUGGUU	689	AACCAUCA CUGAUGAG X CGAA IGAACUUC	2464
1410	AAGUUC C U GAUGGUU	690	AAACCAUC CUGAUGAG X CGAA IGGAACU	2465
1420	AUGGUUUC U GGCAGGA	691	UCCUAGCC CUGAUGAG X CGAA IAAACCAU	2466
1424	UUUCUGGC U AGGAGAGC	692	GCUCUCCU CUGAUGAG X CGAA ICCAGAAA	2467
1433	AGGAGAGC A GCUGGUGU	693	ACACCAGC CUGAUGAG X CGAA ICUCUCCU	2468
1436	AGAGCAGC U GGUGUGCU	694	AGCACACC CUGAUGAG X CGAA ICUGCUCU	2469
1444	UGGUGUGC U GGCAAGCA	695	UGCUUGCC CUGAUGAG X CGAA ICACACCA	2470
1448	GUGCUGGC A AGCAGGCA	696	UGCCUGCU CUGAUGAG X CGAA ICCAGCAC	2471
1452	UGGCAAGC A GGCACCAC	697	GUGGUGCC CUGAUGAG X CGAA ICUUGCCA	2472
1456	AAGCAGGC A CCACCCU	698	AGGGUGG CUGAUGAG X CGAA ICCUGCUU	2473
1458	GCAGGCAC C ACCCUUG	699	CAAGGGU CUGAUGAG X CGAA IUGCCUGC	2474
1459	CAGGCACC A CCCUUGG	700	CCAAGGG CUGAUGAG X CGAA IGUGCCUG	2475
1461	GGCACCAC C CCUUGGA	701	UUCCAAG CUGAUGAG X CGAA IUGGUGCC	2476
1462	GCACCAC C CUUGAAC	702	GUUCCA CUGAUGAG X CGAA IGUGGUGC	2477
1463	CACACCC C UUGGAACA	703	GUUCCAA CUGAUGAG X CGAA IGGUGGUG	2478
1464	ACCACCC C UUGAACAU	704	AUGUCCA CUGAUGAG X CGAA IGGUGGU	2479
1471	CUUGGAAC A UUUUCCA	705	UGGAAAA CUGAUGAG X CGAA IUUCCAAG	2480
1477	ACAUUUUC C CAGUCAUC	706	GAUGACUG CUGAUGAG X CGAA IAAAAUGU	2481
1478	CAUUUUC C AGUCAUCU	707	AGAUGACU CUGAUGAG X CGAA IGAAAAUG	2482
1479	AUUUUC C GUCAUCUC	708	GAGAUGAC CUGAUGAG X CGAA IGGAAAAU	2483
1483	UCCAGUC A UCUCACUC	709	GAGUGAGA CUGAUGAG X CGAA IACUGGGA	2484
1486	CAGUCAUC U CACUCUAC	710	GUAGAGUG CUGAUGAG X CGAA IAUGACUG	2485
1488	GUCAUCUC A CUCUACCU	711	AGGUAGAG CUGAUGAG X CGAA IAGAUGAC	2486
1490	CAUCUCAC U CUACCUAA	712	UUAGGUAG CUGAUGAG X CGAA IUGAGAUG	2487
1492	UCUCACUC U ACCUAAUG	713	CAUUAGGU CUGAUGAG X CGAA IAGUGAGA	2488
1495	CACUCUAC C UAAUGGGU	714	ACCCAUA CUGAUGAG X CGAA IUAGAGUG	2489
1496	ACUCUAC C AAUGGGUG	715	CACCAU CUGAUGAG X CGAA IUGAGAGU	2490
1512	GAGGUUAC C AACCAGUC	716	GACUGGU CUGAUGAG X CGAA IUAACCUC	2491
1513	AGGUUAC C ACCAGUCC	717	GGACUGGU CUGAUGAG X CGAA IGUAACCU	2492
1516	UUACCAAC C AGUCCUUC	718	GAAGGACU CUGAUGAG X CGAA IUUGGUAA	2493
1517	UACCAAC C GUCCUUC	719	GGAAGGAC CUGAUGAG X CGAA IGUGGUAA	2494
1521	AACCAGUC C UCCGCAU	720	AUGCGGA CUGAUGAG X CGAA IACUGGUU	2495
1522	ACCAGUCC U UCCGCAUC	721	GAUGCGGA CUGAUGAG X CGAA IGACUGGU	2496
1525	AGUCCUUC C GCAUCACC	722	GGUGAUGC CUGAUGAG X CGAA IAAGGACU	2497
1528	CCUCCGC A UCACCAUC	723	GAUGGUGA CUGAUGAG X CGAA ICGAAGG	2498
1531	UCCGCAUC A CCAUCCU	724	AAGGAUGG CUGAUGAG X CGAA IAUGCGGA	2499
1533	CGCAUCAC C AUCCUUC	725	GGAAGGAU CUGAUGAG X CGAA IUGAUGCG	2500
1534	GCAUCACC A UCCUCCG	726	CGGAAGGA CUGAUGAG X CGAA IGUGAUGC	2501
1537	UCACCAUC C UCCGCGAG	727	CUGCGGA CUGAUGAG X CGAA IAUGGUGA	2502
1538	CACCAUCC U UCCGCGAG	728	GCUGCGGA CUGAUGAG X CGAA IGAUGGUG	2503
1541	CAUCCUUC C GCAGCAAU	729	AUUGCUGC CUGAUGAG X CGAA IAAGGAUG	2504
1544	CCUCCGC A GCAAUACC	730	GGUAUUGC CUGAUGAG X CGAA ICGAAGG	2505
1547	UCCGCGAG C AUACCUUC	731	GCAGGUAU CUGAUGAG X CGAA ICUGCGGA	2506
1552	GCAAUAC C UGCGGCA	732	UGCGGCA CUGAUGAG X CGAA IUUAUUGCU	2507
1553	GCAAUACC U GCGGCCAG	733	CUGCGGC CUGAUGAG X CGAA IGUAUUGC	2508
1559	CCUGCGGC C AGUGGAAG	734	CUUCCACU CUGAUGAG X CGAA ICCGCGAG	2509
1560	CUGCGGC A GUGGAAGA	735	UCUCCAC CUGAUGAG X CGAA IGCCGCGAG	2510
1575	GAUGUGGC C ACGUCCCA	736	UGGACGU CUGAUGAG X CGAA ICCACAUC	2511
1576	AUGUGGC A CGUCCCA	737	UUGGACG CUGAUGAG X CGAA IGCCACAUC	2512
1581	GCCACGUC C CAAGACGA	738	UCGUCUUG CUGAUGAG X CGAA IACGUGGC	2513
1582	CCACGUCC C AAGACGAC	739	GUCGUCU CUGAUGAG X CGAA IGACGUGG	2514

Table 19

1583	CACGUCCC A AGACGACU	740	AGUCGUCU CUGAUGAG X CGAA IGGACGUG	2515
1591	AAGACGAC U GUUACAAG	741	CUUGUAAC CUGAUGAG X CGAA IUCGUCU	2516
1597	ACUGUAC A AGUUGCC	742	GGCAAACU CUGAUGAG X CGAA IUAACAGU	2517
1605	AAGUUGC C AUCUCACA	743	UGUGAGAU CUGAUGAG X CGAA ICAAACU	2518
1606	AGUUGCC A UCUCACAG	744	CUGUGAGA CUGAUGAG X CGAA IGCAAACU	2519
1609	UUGCCAUC U CACAGUCA	745	UGACUGUG CUGAUGAG X CGAA IAUGGCAA	2520
1611	GCCAUCUC A CAGUCAUC	746	GAUGACUG CUGAUGAG X CGAA IAGAUGGC	2521
1613	CAUCUCAC A GUCAUCCA	747	UGGAUGAC CUGAUGAG X CGAA IUGAGAUG	2522
1617	UCACAGUC A UCCACGGG	748	CCCUGGA CUGAUGAG X CGAA IACUGUGA	2523
1620	CAGUCAUC C ACGGGCAC	749	GUGCCCGU CUGAUGAG X CGAA IAUGACUG	2524
1621	AGUCAUCC A CGGGCACU	750	AGUGCCCG CUGAUGAG X CGAA IGAUGACU	2525
1627	CCACGGG C A CUGUUAUG	751	CAUAACAG CUGAUGAG X CGAA ICCCGUGG	2526
1629	ACGGGCAC U GUUAUGGG	752	CCCAUAC CUGAUGAG X CGAA IUGCCCGU	2527
1641	AUGGGAGC U GUUAUCAU	753	AUGAUAAC CUGAUGAG X CGAA ICUCCCAU	2528
1648	CUGUUAUC A UGGAGGGC	754	GCCCUCCA CUGAUGAG X CGAA IAUAACAG	2529
1657	UGGAGGGC U UCUACGUU	755	AACGUAGA CUGAUGAG X CGAA ICCCUCCA	2530
1660	AGGGCUUC U ACGUUGUC	756	GACAACU CUGAUGAG X CGAA IAAGCCCU	2531
1669	ACGUUGUC U UUGAUCGG	757	CCGAUCAA CUGAUGAG X CGAA IACAACGU	2532
1680	GAUCGGGC C CGAAACG	758	CGUUUUCG CUGAUGAG X CGAA ICCCGAUC	2533
1681	AUCGGGCC C GAAAACGA	759	UCGUUUC CUGAUGAG X CGAA IGCCGAU	2534
1696	GAAUUGGC U UUGCUGUC	760	GACAGCAA CUGAUGAG X CGAA ICCAAUUC	2535
1701	GGCUUUGC U GUCAGCGC	761	GCGCUGAC CUGAUGAG X CGAA ICAAAGCC	2536
1705	UUGCUGUC A GCGCUUGC	762	GCAAGCGC CUGAUGAG X CGAA IACAGCAA	2537
1710	GUCAGCGC U UGCCAUGU	763	ACAUGGCA CUGAUGAG X CGAA ICGCUGAC	2538
1714	GCGCUUGC C AUGUGCAC	764	GUGCACA CUGAUGAG X CGAA ICAAGCGC	2539
1715	CGCUUGCC A UGUGCACG	765	CGUGCACA CUGAUGAG X CGAA IGCAAGCG	2540
1721	CCAUGUGC A CGAUGAGU	766	ACUCAUCG CUGAUGAG X CGAA ICACAUGG	2541
1732	AUGAGUUC A GGACGGCA	767	UGCCGUCC CUGAUGAG X CGAA IAACUCAU	2542
1740	AGGACGGC A GCGUGGA	768	UCCACCGC CUGAUGAG X CGAA ICCGUCCU	2543
1753	UGGAAGGC C CUUUUGUC	769	GACAAAAG CUGAUGAG X CGAA ICCUCCA	2544
1754	GGAAGGCC C UUUUGUCA	770	UGACAAA CUGAUGAG X CGAA IGCCUCC	2545
1755	GAAGGCC U UUGUCAC	771	GUGACAAA CUGAUGAG X CGAA IGGCCUUC	2546
1762	CUUUUGUC A CCUUGGAC	772	GUCCAAGG CUGAUGAG X CGAA IACAAAAG	2547
1764	UUUGUCAC C UUGGACAU	773	AUGUCAA CUGAUGAG X CGAA IUAGACAA	2548
1765	UUGUCACC U UGGACAUG	774	CAUGUCCA CUGAUGAG X CGAA IGUGACAA	2549
1771	CCUUGGAC A UGGAAGAC	775	GUCUCCA CUGAUGAG X CGAA IUCCAAGG	2550
1780	UGGAAGAC U GUGGCUAC	776	GUAGCCAC CUGAUGAG X CGAA IUCUCCA	2551
1786	ACUGUGGC U ACAACAUU	777	AAUGUUGU CUGAUGAG X CGAA ICCACAGU	2552
1789	GUGGCUAC A ACAUCCA	778	UGGAAUGU CUGAUGAG X CGAA IUAGCCAC	2553
1792	GCUACAAC A UUCCACAG	779	CUGUGGAA CUGAUGAG X CGAA IUUGUAGC	2554
1796	CAACAUUC C ACAGACAG	780	CUGUCUGU CUGAUGAG X CGAA IAAUGUUG	2555
1797	AACAUUCC A CAGACAGA	781	UCUGUCUG CUGAUGAG X CGAA IGAUGUU	2556
1799	CAUCCAC A GACAGAUG	782	CAUCUGUC CUGAUGAG X CGAA IUGGAAUG	2557
1803	CCACAGAC A GAUGAGUC	783	GACUCAUC CUGAUGAG X CGAA IUCUGUGG	2558
1812	GAGAGUC A ACCCUCAU	784	AUGAGGGU CUGAUGAG X CGAA IACUCAUC	2559
1815	GAGUCAAC C CUCAUGAC	785	GUCAUGAG CUGAUGAG X CGAA IUUGACUC	2560
1816	AGUCAACC C UCAUGACC	786	GGUCAUGA CUGAUGAG X CGAA IGUGACU	2561
1817	GUCAACCC U CAUGACCA	787	UGGUCAUG CUGAUGAG X CGAA IGGUUGAC	2562
1819	CAACCCUC A UGACCAUA	788	UAUGGUCA CUGAUGAG X CGAA IAGGUUG	2563
1824	CUCAUGAC C AUAGCCUA	789	UAGGCUAU CUGAUGAG X CGAA IUCAUGAG	2564
1825	UCAUGACC A UAGCCUAU	790	AUAGGCUA CUGAUGAG X CGAA IGUCAUGA	2565

Table 19

1830	ACCAUAGC C UAUGUCAU	791	AUGACAU CUGAUGAG X CGAA ICUAUGGU	2566
1831	CCAUGACC U AUGUCAUG	792	CAUGACAU CUGAUGAG X CGAA IGCUAUGG	2567
1837	CCUAUGUC A UGGCUGCC	793	GGCAGCCA CUGAUGAG X CGAA IACAUAGG	2568
1842	GUCAUGGC U GCCAUCUG	794	CAGAUGGC CUGAUGAG X CGAA ICCAUGAC	2569
1845	AUGGCUGC C AUCUGCGC	795	GCGCAGAU CUGAUGAG X CGAA ICAGCCA	2570
1846	UGGCUGCC A UCUGCGCC	796	GGCGCAGA CUGAUGAG X CGAA IGCAGCCA	2571
1849	CUGCCAUC U GCGCCUC	797	GAGGGCGC CUGAUGAG X CGAA IAUGGCAG	2572
1854	AUCUGCGC C CUCUUCAU	798	AUGAAGAG CUGAUGAG X CGAA ICGCAGAU	2573
1855	UCUGCGCC C UCUCUAUG	799	CAUGAAGA CUGAUGAG X CGAA ICGCAGAU	2574
1856	CUGCGCCC U CUUCAUGC	800	GCAUGAAG CUGAUGAG X CGAA IGGCGCAG	2575
1858	GCGCCUC U UCAUGCUG	801	CAGCAUGA CUGAUGAG X CGAA IAGGGCGC	2576
1861	CCCUCUUC A UGCUGCCA	802	UGGCAGCA CUGAUGAG X CGAA IAAGAGGG	2577
1865	CUUCAUGC U GCCACUCU	803	AGAGUGGC CUGAUGAG X CGAA ICAUGAAG	2578
1868	CAUGCUGC C ACUCUGCC	804	GGCAGAGU CUGAUGAG X CGAA ICAGCAUG	2579
1869	AUGCUGCC A CUCUGCCU	805	AGGCAGAG CUGAUGAG X CGAA IGCAGCAU	2580
1871	GCUGCCAC U CUGCCUCA	806	UGAGGCAG CUGAUGAG X CGAA IUGGCAGC	2581
1873	UGCCACUC U GCCUCAUG	807	CAUGAGGC CUGAUGAG X CGAA IAGUGGCA	2582
1876	CACUCUGC C UCAUGGUG	808	CACCAUGA CUGAUGAG X CGAA ICAGAGUG	2583
1877	ACUCUGCC U CAUGGUGU	809	ACACCAUG CUGAUGAG X CGAA IGCAGAGU	2584
1879	UCUGCCUC A UGGUGUGU	810	ACACACCA CUGAUGAG X CGAA IAGGCAGA	2585
1889	GGUGUGUC A GUGGCGCU	811	AGCGCCAC CUGAUGAG X CGAA IACACACC	2586
1897	AGUGGCGC U GCCUCCGC	812	GCGGAGGC CUGAUGAG X CGAA ICGCCACU	2587
1900	GGCGCUGC C UCCGUGGC	813	GCAGCGGA CUGAUGAG X CGAA ICAGCGCC	2588
1901	GCGCUGCC U CCGUGGCC	814	GGCAGCGG CUGAUGAG X CGAA IGCAGCGC	2589
1903	GCUGCCUC C GCUGCCUG	815	CAGGCAGC CUGAUGAG X CGAA IAGGCAGC	2590
1906	GCCUCCGC U GCCUGCGC	816	GCGCAGGC CUGAUGAG X CGAA ICGGAGGC	2591
1909	UCCGUGC C UGCGCCAG	817	CUGGCGCA CUGAUGAG X CGAA ICAGCGGA	2592
1910	CCGUGCC U GCGCCAGC	818	GCUGGCGC CUGAUGAG X CGAA IGCAGCGG	2593
1915	GCCUGCGC C AGCAGCAU	819	AUGCUGCU CUGAUGAG X CGAA ICGCAGGC	2594
1916	CCUGCGCC A GCAGCAUG	820	CAUGCUGC CUGAUGAG X CGAA ICGCAGG	2595
1919	GCGCCAGC A GCAUGAUG	821	CAUCAUGC CUGAUGAG X CGAA ICUGCGC	2596
1922	CCAGCAGC A UGAUGACU	822	AGUCAUCA CUGAUGAG X CGAA ICUGCUGG	2597
1930	AUGAUGAC U UUGCUGAU	823	AUCAGCAA CUGAUGAG X CGAA IUCAUCAU	2598
1935	GACUUGC U GAUGACAU	824	AUGUCAUC CUGAUGAG X CGAA ICAAAGUC	2599
1942	CUGAUGAC A UCUCUCCUG	825	CAGGGAGA CUGAUGAG X CGAA IUCAUCAG	2600
1945	AUGACAUC U CCCUGCUG	826	CAGCAGGG CUGAUGAG X CGAA IAUGUCAU	2601
1947	GACAUCUC C CUGCUGAA	827	UUCAGCAG CUGAUGAG X CGAA IAGAUGUC	2602
1948	ACAUCUCC C UGCUGAAG	828	CUUCAGCA CUGAUGAG X CGAA IGAGAUGU	2603
1949	CAUCUCCC U GCUGAAGU	829	ACUUCAGC CUGAUGAG X CGAA IGGAGAUG	2604
1952	CUCCUGC U GAAGUGAG	830	CUCACUUC CUGAUGAG X CGAA ICAGGGAG	2605
1966	GAGGAGGC C CAUGGGCA	831	UGCCCAUG CUGAUGAG X CGAA ICCUCCUC	2606
1967	AGGAGGCC C AUGGGCAG	832	CUGCCCAU CUGAUGAG X CGAA IGCCUCCU	2607
1968	GGAGGCCC A UGGGCAGA	833	UCUGCCCA CUGAUGAG X CGAA IGGCCUCC	2608
1974	CCAUGGGC A GAAGAUAG	834	CUAUCUCC CUGAUGAG X CGAA ICCCAUGG	2609
1989	AGAGAUUC C CCUGGACC	835	GGUCCAG CUGAUGAG X CGAA IAAUCUCU	2610
1990	GAGAUUCC C CUGGACCA	836	UGGUCCAG CUGAUGAG X CGAA IGAAUCUC	2611
1991	AGAUUCCC C UGACCAC	837	GUGGUCCA CUGAUGAG X CGAA IGGAAUCU	2612
1992	GAUUCCCC U GGACCACA	838	UGUGGUCC CUGAUGAG X CGAA IUCCAGG	2613
1997	CCUGGAC C ACACUCC	839	GGAGGUGU CUGAUGAG X CGAA IUCCAGG	2614
1998	CCUGGACC A CACCUCCG	840	CGGAGGUG CUGAUGAG X CGAA IGUCAGG	2615
2000	UGGACCAC A CCUCCGUG	841	CACGGAGG CUGAUGAG X CGAA IUGGUCCA	2616

Table 19

2002	GACCACAC C UCCGUGGU	842	ACCACGGA CUGAUGAG X CGAA IUGUGGUC	2617
2003	ACCACACC U CCGUGGUU	843	AACCACGG CUGAUGAG X CGAA IGUGUGGU	2618
2005	CACACCUC C GUGGUUCA	844	UGAACCAC CUGAUGAG X CGAA IAGGUGUG	2619
2013	CGUGGUUC A CUUUGGUC	845	GACCAAAG CUGAUGAG X CGAA IAACCACG	2620
2015	UGGUUCAC U UUGGUCAC	846	GUGACCAA CUGAUGAG X CGAA IUGAACCA	2621
2022	CUUUGGUC A CAAGUAGG	847	CCUACUUG CUGAUGAG X CGAA IACCAAAG	2622
2024	UUGGUCAC A AGUAGGAG	848	CUCCUACU CUGAUGAG X CGAA IUGACCAA	2623
2035	UAGGAGAC A CAGAUGGC	849	GCCAUCUG CUGAUGAG X CGAA IUCUCCUA	2624
2037	GGAGACAC A GAUGGCAC	850	GUGCCAUC CUGAUGAG X CGAA IUGUCUCC	2625
2044	CAGAUGGC A CCUGUGGC	851	GCCACAGG CUGAUGAG X CGAA ICCAUCUG	2626
2046	GAUGGCAC C UGUGGCCA	852	UGGCCACA CUGAUGAG X CGAA IUGCCAUC	2627
2047	AUGGCACC U GUGGCCAG	853	CUGGCCAC CUGAUGAG X CGAA IGUGCCAU	2628
2053	CCUGUGGC C AGAGCACC	854	GGUGCUCU CUGAUGAG X CGAA ICCACAGG	2629
2054	CUGUGGCC A GAGCACC	855	AGGUGCUC CUGAUGAG X CGAA IGCCACAG	2630
2059	GCCAGAGC A CCUCAGGA	856	UCCUGAGG CUGAUGAG X CGAA ICUCUGGC	2631
2061	CAGAGCAC C UCAGGACC	857	GGUCCUGA CUGAUGAG X CGAA IUGCUCUG	2632
2062	AGAGCACC U CAGGACCC	858	GGGUCCUG CUGAUGAG X CGAA IGUGCUCU	2633
2064	AGCACCUC A GGACCCUC	859	GAGGGUCC CUGAUGAG X CGAA IAGGUGCU	2634
2069	CUCAGGAC C CUCCCCAC	860	GUGGGGAG CUGAUGAG X CGAA IUCCUGAG	2635
2070	UCAGGACC C UCCCCACC	861	GGUGGGGA CUGAUGAG X CGAA IGUCCUGA	2636
2071	CAGGACCC U CCCCACCC	862	GGGUGGGG CUGAUGAG X CGAA IGGUCCUG	2637
2073	GGACCCUC C CCACCCAC	863	GUGGGUGG CUGAUGAG X CGAA IAGGGUCC	2638
2074	GACCCUCC C CACCCACC	864	GGUGGGUG CUGAUGAG X CGAA IGAGGGUC	2639
2075	ACCCUCCC C ACCCACCA	865	UGSUGGGU CUGAUGAG X CGAA IGGAGGGU	2640
2076	CCCUCCCC A CCCACCAA	866	UUGGUGGG CUGAUGAG X CGAA IGGGAGGG	2641
2078	CUCCCCAC C CACCAAAU	867	AUUUGGUG CUGAUGAG X CGAA IUGGGGAG	2642
2079	UCCCCACC C ACCAAAU	868	CAUUUGGU CUGAUGAG X CGAA IGUGGGGA	2643
2080	CCCCACCC C CCAAUUGC	869	GCAUUUGG CUGAUGAG X CGAA IGGUGGGG	2644
2082	CCACCCAC C AAAUGCCU	870	AGGCAUUU CUGAUGAG X CGAA IUUGGUGG	2645
2083	CACCCACC A AAUGCCUC	871	GAGGCAUU CUGAUGAG X CGAA IGUGGGUG	2646
2089	CCAAUUGC C UCUGCCUU	872	AAGGCAGA CUGAUGAG X CGAA ICAUUUGG	2647
2090	CAAUUGCC U CUGCCUUG	873	CAAGGCAG CUGAUGAG X CGAA IGCAUUUG	2648
2092	AAUGCCUC U GCCUUGAU	874	AUCAAGGC CUGAUGAG X CGAA IAGGCAUU	2649
2095	GCCUCUGC C UUGAUGGA	875	UCCAUCAA CUGAUGAG X CGAA ICAGAGGC	2650
2096	CCUCUGCC U UGAUGGAG	876	CUCCAUCA CUGAUGAG X CGAA IGCAGAGG	2651
2116	GAAAAGGC U GGCAAGGU	877	ACCUUGCC CUGAUGAG X CGAA ICCUUUUC	2652
2120	AGGCUGGC A AGGUGGGU	878	ACCCACCU CUGAUGAG X CGAA ICCAGCCU	2653
2131	GUGGGUUC C AGGGACUG	879	CAGUCCCU CUGAUGAG X CGAA IAACCCAC	2654
2132	UGGGUUC C A GGGACUGU	880	ACAGUCCC CUGAUGAG X CGAA IGAACCCA	2655
2138	CCAGGGAC U GUACCUGU	881	ACAGGUAC CUGAUGAG X CGAA IUCCUGG	2656
2143	GACUGUAC C UGUAGGAA	882	UUCCUACA CUGAUGAG X CGAA IUACAGUC	2657
2144	ACUGUACC U GUAGGAAA	883	UUUCCUAC CUGAUGAG X CGAA IGUACAGU	2658
2154	UAGGAAAC A GAAAAGAG	884	CUCUUUUC CUGAUGAG X CGAA IUUUCCUA	2659
2174	AAAGAAGC A CUCUGCUG	885	CAGCAGAG CUGAUGAG X CGAA ICUUUUUU	2660
2176	AGAAGCAC U CUGUGGC	886	GCCAGCAG CUGAUGAG X CGAA IUGCUUCU	2661
2178	AAGCACUC U GCUGCGG	887	CCGCCAGC CUGAUGAG X CGAA IAGUGCUU	2662
2181	CACUCUGC U GCGGGGAA	888	UUCCCGCC CUGAUGAG X CGAA ICAGAGUG	2663
2193	GGGAAUAC U CUUGGUCA	889	UGACCAAG CUGAUGAG X CGAA IUUUUUCC	2664
2195	GAAUACUC U UGGUCACC	890	GGUGACCA CUGAUGAG X CGAA IAGUAUUC	2665
2201	UCUUGGUC A CCUCAAAU	891	AUUUGAGG CUGAUGAG X CGAA IACCAAGA	2666
2203	UUGGUCAC C UCAAAUUU	892	AAAUUGA CUGAUGAG X CGAA IUGACCAA	2667

Table 19

2204	UGGUCACC U CAAAUUUA	893	UAAAUUUG CUGAUGAG X CGAA IGUGACCA	2668
2206	GUCACCUC A AAUUUAAG	894	CUUAAAUU CUGAUGAG X CGAA IAGGUGAC	2669
2226	GGAAAUUC U GCUGCUUG	895	CAAGCAGC CUGAUGAG X CGAA IAAUUUCC	2670
2229	AAUUCUGC U GCUGGAAA	896	UUUCAAGC CUGAUGAG X CGAA ICAGAAUU	2671
2232	UCUGCUGC U UGAAACUU	897	AAGUUUCA CUGAUGAG X CGAA ICAGCAGA	2672
2239	CUUGAAAC U UCAGCCCU	898	AGGGCUGA CUGAUGAG X CGAA IUUUCAAG	2673
2242	GAAACUUC A GCCCUGAA	899	UUCAGGGC CUGAUGAG X CGAA IAAGUUUC	2674
2245	ACUUCAGC C CUGAACCU	900	AGGUUCAG CUGAUGAG X CGAA ICUGAAGU	2675
2246	CUUCAGCC C UGAACCUU	901	AAGGUUCA CUGAUGAG X CGAA IGCUGAAG	2676
2247	UUCAGCCC U GAACCUUU	902	AAAGGUUC CUGAUGAG X CGAA IGGCUGAA	2677
2252	CCUGAAC C UUUGUCCA	903	UGGACAAA CUGAUGAG X CGAA IUUCAGGG	2678
2253	CCUGAAC U UUGUCCAC	904	GUGGACAA CUGAUGAG X CGAA IGUUCAGG	2679
2259	CCUUGUC C ACCAUUCC	905	GGAAUGGU CUGAUGAG X CGAA IACAAAGG	2680
2260	CUUUGUCC A CCAUUCUU	906	AGGAAUGG CUGAUGAG X CGAA IGACAAAG	2681
2262	UUGUCCAC C AUUCCUUU	907	AAAGGAAU CUGAUGAG X CGAA IUGGACAA	2682
2263	UGUCCACC A UJCCUUUA	908	UAAAGGAA CUGAUGAG X CGAA IGUGGACA	2683
2267	CACCAUUC C UUUAAAUU	909	AAUUUAAA CUGAUGAG X CGAA IAAUGGUG	2684
2268	ACCAUUC U UUAUUUUC	910	GAAUUUAA CUGAUGAG X CGAA IGAUUGGU	2685
2277	UUAAAUUC U CCAACCCA	911	UGGGUUGG CUGAUGAG X CGAA IAAUUUAA	2686
2279	AAAUUCUC C AACCCAAA	912	UUUGGGUU CUGAUGAG X CGAA IAGAAUUU	2687
2280	AAUUCUCC A ACCCAAAG	913	CUUUGGGU CUGAUGAG X CGAA IGACAAUU	2688
2283	UCUCCAAC C CAAAGUUA	914	AUACUUUG CUGAUGAG X CGAA IUUGGAGA	2689
2284	CUCCAACC C AAAGUAUU	915	AAUACUUU CUGAUGAG X CGAA IGUUGGAG	2690
2285	UCCAACCC A AAGUAUUC	916	GAAUACUU CUGAUGAG X CGAA IGGUUGGA	2691
2294	AAGUAUUC U UCUUUUCU	917	AGAAAAGA CUGAUGAG X CGAA IAAUACUU	2692
2297	UAUUCUUC U UUUCUUAG	918	CUAAGAAA CUGAUGAG X CGAA IAAAGAAU	2693
2302	UUCUUUUC U UAGUUUCA	919	UGAAACUA CUGAUGAG X CGAA IAAAAGAA	2694
2310	UUAGUUUC A GAAGUACU	920	AGUACUUC CUGAUGAG X CGAA IAAACUAA	2695
2318	AGAAGUAC U GGCAUCAC	921	GUGAUGCC CUGAUGAG X CGAA IUACUUCU	2696
2322	GUACUGGC A UCACACGC	922	GCGUGUGA CUGAUGAG X CGAA ICCAGUAC	2697
2325	CUGGCAUC A CACGCAGG	923	CCUGCGUG CUGAUGAG X CGAA IAUGCCAG	2698
2327	GGCAUCAC A CGCAGGUU	924	AACCUGCG CUGAUGAG X CGAA IUUGAUGCC	2699
2331	UCACACGC A GGUUACCU	925	AGGUAAAC CUGAUGAG X CGAA ICGUGUGA	2700
2338	CAGGUUAC C UUGGCGUG	926	CACGCCAA CUGAUGAG X CGAA IUAACCUG	2701
2339	AGGUUACC U UGGCGUGU	927	ACACGCCA CUGAUGAG X CGAA IGUAACCU	2702
2351	CGUGUGUC C CUGUGGUA	928	UACCACAG CUGAUGAG X CGAA IACACACG	2703
2352	GUGUGUCC C UGUGGUAC	929	GUACCACA CUGAUGAG X CGAA IGACACAC	2704
2353	UGUGUCCC U GUGGUACC	930	GGUACCAC CUGAUGAG X CGAA IGGACACA	2705
2361	UGUGGUAC C CUGGCAGA	931	UCUGCCAG CUGAUGAG X CGAA IUACCACA	2706
2362	GUGGUUAC C UGGCAGAG	932	CUCUGCCA CUGAUGAG X CGAA IGUACCAC	2707
2363	UGGUACCC U GGCAGAGA	933	UCUCUGCC CUGAUGAG X CGAA IGGUACCA	2708
2367	ACCCUGGC A GAGAAGAG	934	CUCUUCUC CUGAUGAG X CGAA ICCAGGGU	2709
2378	GAAGAGAC C AAGCUUGU	935	ACAAGCUU CUGAUGAG X CGAA IUCUCUUC	2710
2379	AAGAGACC A AGCUUGUU	936	AACAAGCU CUGAUGAG X CGAA IGUCUCUU	2711
2383	GACCAAGC U UGUUCCCC	937	GGGAAACA CUGAUGAG X CGAA ICUUGGUC	2712
2390	CUUGUUUC C CUGCUGGC	938	GCCAGCAG CUGAUGAG X CGAA IAAACAAG	2713
2391	UUGUUUCC C UGCUGGCC	939	GGCCAGCA CUGAUGAG X CGAA IGAAACAA	2714
2392	UGUUUCCC U GCUGGCCA	940	UGGCCAGC CUGAUGAG X CGAA IGGAAACA	2715
2395	UUCCCUGC U GGCCAAAG	941	CUUUGGCC CUGAUGAG X CGAA ICAGGGAA	2716
2399	CUGCUGGC C AAAGUCAG	942	CUGACUUU CUGAUGAG X CGAA ICCAGCAG	2717
2400	UGCUGGCC A AAGUCAGU	943	ACUGACUU CUGAUGAG X CGAA IGCCAGCA	2718

Table 19

2406	CCAAAGUC A GUAGGAGA	944	UCUCCUAC CUGAUGAG X CGAA IACUUUGG	2719
2421	GAGGAUGC A CAGUUUGC	945	GCAAACUG CUGAUGAG X CGAA ICAUCCUC	2720
2423	GGAUGCAC A GUUUGCUA	946	UAGCAAAC CUGAUGAG X CGAA IUGCAUCC	2721
2430	CAGUUUGC U AUUUGCUU	947	AAGCAAU CUGAUGAG X CGAA ICAAACUG	2722
2437	CUAUUUGC U UUAGAGAC	948	GUCUCUAA CUGAUGAG X CGAA ICAAUAG	2723
2446	UUAGAGAC A GGGACUGU	949	ACAGUCCC CUGAUGAG X CGAA IUCUCUAA	2724
2452	ACAGGGAC U GUUAAAC	950	GUUUUAC CUGAUGAG X CGAA IUCCCUGU	2725
2461	GUUAAAC A AGCCUAAAC	951	GUUAGGCU CUGAUGAG X CGAA IUUUUAC	2726
2465	AAACAAGC C UAACAUUG	952	CAAUGUUA CUGAUGAG X CGAA ICUUGUUU	2727
2466	AACAAGCC U AACAUUGG	953	CCAAUGUU CUGAUGAG X CGAA IGCUUGUU	2728
2470	AGCCUAAAC A UUGGUGCA	954	UGCACCAA CUGAUGAG X CGAA IUUAGGCU	2729
2478	AUUGGUGC A AAGAUUGC	955	GCAAUCUU CUGAUGAG X CGAA ICACCAU	2730
2487	AAGAUUGC C UCUUGAAU	956	AUUCAAGA CUGAUGAG X CGAA ICAAUCUU	2731
2488	AGAUUGCC U CUUGAAUU	957	AAUUAAG CUGAUGAG X CGAA IGCAAUCU	2732
2490	AUUGCCUC U UGAAUUA	958	UUAAUUA CUGAUGAG X CGAA IAGGCAU	2733
2509	AAAAAAC U AGAAAAA	959	UUUUUUCU CUGAUGAG X CGAA IUUUUUU	2734

Input Sequence = AF190725. Cut Site = G/.

Stem Length = 8. Core Sequence = CUGAUGAG X CGAA (X = GCCGUUAGGC or other stem II)

AF190725 (Homo sapiens beta-site APP cleaving enzyme (BACE) mRNA; 2526 bp)

Table 20

Table 20: Human BACE G-cleaver Ribozyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
11	ACGCGUCC G CAGCCCGC	960	GCGGGCUG UGAUG GCAUGCACUAUGC GCG GGACGCGU	2735
18	CGCAGCCC G CCGGGGAG	961	CUCCCGGG UGAUG GCAUGCACUAUGC GCG GGGCUGGG	2736
29	CGGGAGCU G CGAGCCGC	962	GCGGCUCG UGAUG GCAUGCACUAUGC GCG AGCUCCCG	2737
31	GGAGCUCC G AGCCGCGA	963	UCGCGGCU UGAUG GCAUGCACUAUGC GCG GCAGCUCC	2738
36	UGCGAGCC G CGAGCUGG	964	CCAGCUCG UGAUG GCAUGCACUAUGC GCG GGCUCGGA	2739
38	CGAGCCGC G AGCUGGAU	965	AUCCAGCU UGAUG GCAUGCACUAUGC GCG GCGGCUUG	2740
58	GGUGGCGU G AGCAGCCA	966	UGGCUCUG UGAUG GCAUGCACUAUGC GCG AGGCCACC	2741
69	CAGCCAAC G CAGCCGCA	967	UGCGGCTUG UGAUG GCAUGCACUAUGC GCG GUUGGCUU	2742
75	ACGCAGCC G CAGGAGCC	968	GGCUCCUG UGAUG GCAUGCACUAUGC GCG GGCUGCGU	2743
94	GAGCCCUU G CCCCUGCC	969	GGCAGGGG UGAUG GCAUGCACUAUGC GCG AAGGGCUC	2744
100	UUGCCCCU G CCGCGGCC	970	GGCGCGGG UGAUG GCAUGCACUAUGC GCG AGGGGCAA	2745
104	CCUUGCCC G CGCCGCCG	971	CGCGCGCG UGAUG GCAUGCACUAUGC GCG GGGCAGGG	2746
106	CUGCCCGC G CCGCCGCC	972	GGCGGCGG UGAUG GCAUGCACUAUGC GCG GCGGGCAG	2747
109	CCCGGGCC G CCGCCCGC	973	GCGGGCGG UGAUG GCAUGCACUAUGC GCG GCGGCGGG	2748
112	GCGCGGCC G CCGCGCGG	974	CCGCGCGG UGAUG GCAUGCACUAUGC GCG GCGGCGCG	2749
116	CGCGGCCG G CCGGGGGG	975	CCCCCGGG UGAUG GCAUGCACUAUGC GCG GGGCGGCG	2750
137	GGGAGCC G CCACCGGC	976	GCGGUGGG UGAUG GCAUGCACUAUGC GCG GGCUCUCC	2751
148	ACCGGCCG G CCAUGCCC	977	GCGCAUGG UGAUG GCAUGCACUAUGC GCG GGGCCCGU	2752
153	CCGCGCAU G CCGGCCCC	978	GGGCGCGG UGAUG GCAUGCACUAUGC GCG AUGGCGGG	2753
157	CCAUGCCC G CCCCUCGC	979	GGGAGGGG UGAUG GCAUGCACUAUGC GCG GGGCAUGG	2754
172	CCAGCCCC G CCGGGAGC	980	GCUCGCCG UGAUG GCAUGCACUAUGC GCG GGGGCUUG	2755
183	GGGAGCCC G CGCCCGCU	981	AGCGGGCG UGAUG GCAUGCACUAUGC GCG GGGCUCCC	2756
185	GAGCCCGC G CCGGCUUG	982	GCAGCGGG UGAUG GCAUGCACUAUGC GCG GCGGGCUC	2757
189	CCGCGCCC G CUGCCCGAG	983	CUGGGCAG UGAUG GCAUGCACUAUGC GCG GGGCGCGG	2758
192	CGCCCGCU G CCGAGGCU	984	AGCCUGGG UGAUG GCAUGCACUAUGC GCG AGCGGGCG	2759
205	GGCUGGCC G CCGCCGUG	985	CACGGCGG UGAUG GCAUGCACUAUGC GCG GGCCAGCC	2760
208	UGGCCGCC G CCGUGCCG	986	CGGCACGG UGAUG GCAUGCACUAUGC GCG GCGGCGCA	2761
213	GCCGCCGU G CCGAUGUA	987	UACAUCGG UGAUG GCAUGCACUAUGC GCG ACGGCGGC	2762
216	GCCGUGCC G AUGUAGCG	988	CGCUACAU UGAUG GCAUGCACUAUGC GCG GGCACGGC	2763
250	UCUCCCGU G CUCCCGUG	989	CACGGGAG UGAUG GCAUGCACUAUGC GCG AGGGGAGA	2764

Table 20

258	GCUCGCCU G CUCUGCGG	990	CCGCAGAG UGAUG GCAUGCACUAUGC GCG ACGGGAGC	2765
263	CGUGCUCU G CGAUCUC	991	GAGAUCCG UGAUG GCAUGCACUAUGC GCG AGAGCACG	2766
276	UCUCCCCU G ACCGUCU	992	AGAGCGGU UGAUG GCAUGCACUAUGC GCG AGGGGAGA	2767
280	CCUGACC G CUCUCCAC	993	GUGGAGAG UGAUG GCAUGCACUAUGC GCG GGUACAGG	2768
320	AGGGCCCU G CAGGCCCU	994	AGGGCCUG UGAUG GCAUGCACUAUGC GCG AGGGCCCU	2769
337	GGGUCUCU G AUGCCCC	995	GGGGCAU UGAUG GCAUGCACUAUGC GCG AGGACGCC	2770
340	GUCCUGAU G CCCCCAAG	996	CUUGGGG UGAUG GCAUGCACUAUGC GCG AUCAGGAC	2771
360	CCUCUCCU G AGRAGCCA	997	UGGCUUCU UGAUG GCAUGCACUAUGC GCG AGGAGAGG	2772
397	GGGCAGGC G CCAGGGAC	998	GUCCUUGG UGAUG GCAUGCACUAUGC GCG GCUUGCCC	2773
420	GGGCCAGU G CGAGCCCA	999	UGGGCUCG UGAUG GCAUGCACUAUGC GCG ACUGGGCC	2774
422	GCCAGUGC G AGCCAGA	1000	UCUGGGU UGAUG GCAUGCACUAUGC GCG GCACUGGC	2775
437	GAGGCCG G AAGGCCG	1001	CCGGCCU UGAUG GCAUGCACUAUGC GCG GGGCCUC	2776
468	CAAGCCCU G CCCUGGU	1002	AGCCAGGG UGAUG GCAUGCACUAUGC GCG AGGGCUG	2777
480	UGGCUCCU G CUGUGGAU	1003	AUCCACAG UGAUG GCAUGCACUAUGC GCG AGGAGCCA	2778
493	GGAUGGC G CGGGAGUG	1004	CACUCCG UGAUG GCAUGCACUAUGC GCG GCCAUCC	2779
501	GCGGAGU G CUGCCUGC	1005	GCAGCAG UGAUG GCAUGCACUAUGC GCG ACUCCCGC	2780
504	GGAGUGCU G CCUGCCCA	1006	UGGCAGG UGAUG GCAUGCACUAUGC GCG AGCACUCC	2781
508	UGCUGCCU G CCCACGGC	1007	GCCUGGG UGAUG GCAUGCACUAUGC GCG AGGCAGCA	2782
537	AUCCGGCU G CCCUGCG	1008	CGCAGGG UGAUG GCAUGCACUAUGC GCG AGCCGGAU	2783
543	CUGCCCCU G CGCAGCGG	1009	CCGUCGCG UGAUG GCAUGCACUAUGC GCG AGGGGCAG	2784
545	GCCCCUGC G CAGCGGCC	1010	GGCCGCG UGAUG GCAUGCACUAUGC GCG GCAGGGGC	2785
562	UGGGGGC G CCCCCUG	1011	CAGGGGG UGAUG GCAUGCACUAUGC GCG GCCCCCCA	2786
576	CUGGGGCU G CGGUCGCC	1012	GGCAGCG UGAUG GCAUGCACUAUGC GCG AGCCCCAG	2787
582	CUGCGGCU G CCCCAGGA	1013	UCCGGGG UGAUG GCAUGCACUAUGC GCG AGCCGCG	2788
595	GGGAGACC G ACGAAGAG	1014	CUCUUGU UGAUG GCAUGCACUAUGC GCG GGUCCUCC	2789
598	AGACCGAC G AAGAGCCC	1015	GGGCUUU UGAUG GCAUGCACUAUGC GCG GUCGGUCU	2790
607	AAGAGCCC G AGGAGCCC	1016	GGGCUUU UGAUG GCAUGCACUAUGC GCG GGGCUCUU	2791
654	GACAACCU G AGGGGCAA	1017	UUGCCCCU UGAUG GCAUGCACUAUGC GCG AGGUUGUC	2792
690	GUGGAGAU G ACCGUGGG	1018	CCCACGU UGAUG GCAUGCACUAUGC GCG AUCUCCAC	2793
708	AGCCCCC G CAGACGCU	1019	AGCUCUG UGAUG GCAUGCACUAUGC GCG GGGGGCU	2794
714	CCGCAGAC G CUCAACAU	1020	AUGUAGAG UGAUG GCAUGCACUAUGC GCG GUCUGCGG	2795
751	GUAAUUU G CAGUGGGU	1021	ACCCACUG UGAUG GCAUGCACUAUGC GCG AAAGUUAC	2796
760	CAGUGGGU G CUGCCCCC	1022	GGGGCAG UGAUG GCAUGCACUAUGC GCG ACCCACUG	2797
763	UGGGUGCU G CCCCCCAC	1023	GUGGGGG UGAUG GCAUGCACUAUGC GCG AGCACCCA	2798

Table 20

780	CCCUUCCU G CAUCGCUA	1024	UACGGAUG UGAUG GCAUGCACUAUGC GCG AGGAAGGG	2799
785	CCUGCAUC G CUACUACC	1025	GGUAGUAG UGAUG GCAUGCACUAUGC GCG GAUGCAGG	2800
843	GUGUAUGU G CCUACAC	1026	GUGUAGGG UGAUG GCAUGCACUAUGC GCG ACAUACAC	2801
883	UGGGCACC G ACCUGGUA	1027	UACCAGGU UGAUG GCAUGCACUAUGC GCG GGUGCCCA	2802
921	GUCACUGU G CGUGCCAA	1028	UUGGCACG UGAUG GCAUGCACUAUGC GCG ACAGUGAC	2803
925	CUGUGCGU G CCAACAUU	1029	AAUUGUGG UGAUG GCAUGCACUAUGC GCG ACGCACAG	2804
934	CCAACAUU G CUGCCAUU	1030	GAUGGCAG UGAUG GCAUGCACUAUGC GCG AAUGUUGG	2805
937	ACAUUGCU G CCAUCACU	1031	AGUGAUGG UGAUG GCAUGCACUAUGC GCG AGCAAUGU	2806
946	CCAUCACU G AUUCAGAC	1032	GUCUGAUU UGAUG GCAUGCACUAUGC GCG AGUGAUGG	2807
1006	UGGCCUUAU G CUGAGAUU	1033	AAUCUCAG UGAUG GCAUGCACUAUGC GCG AUAGGCCA	2808
1009	CCUAUGCU G AGAUUGCC	1034	GGCAUUCU UGAUG GCAUGCACUAUGC GCG AGCAUAGG	2809
1015	CUGAGAUU G CCAGGCCU	1035	AGGCCUGG UGAUG GCAUGCACUAUGC GCG AAUCUCAG	2810
1024	CCAGGCCU G ACACUCC	1036	GGAGUCGU UGAUG GCAUGCACUAUGC GCG AGGCCUUG	2811
1027	GGCCUGAC G ACUCCUG	1037	CAGGGAGU UGAUG GCAUGCACUAUGC GCG GUCAGGCC	2812
1048	CUUUCUUU G ACUCUCUG	1038	CAGAGAGU UGAUG GCAUGCACUAUGC GCG AAAGAAAG	2813
1092	UUCUCCCU G CAGCUUUG	1039	CAAAGCUG UGAUG GCAUGCACUAUGC GCG AGGGAGAA	2814
1105	UUUGUGGU G CUGGCUUC	1040	GAAGCCAG UGAUG GCAUGCACUAUGC GCG ACCACAAA	2815
1129	ACCAGUCU G AAGUGCUG	1041	CAGCACUU UGAUG GCAUGCACUAUGC GCG AGACUGGU	2816
1134	UCUGAAGU G CUGGCCUC	1042	GAGGCCAG UGAUG GCAUGCACUAUGC GCG ACUUCAGA	2817
1158	GGGAGCAU G AUCAUUGG	1043	CCAAUGAU UGAUG GCAUGCACUAUGC GCG AUGCUCCC	2818
1174	GAGGUADC G ACCACUCG	1044	CGAGUGGU UGAUG GCAUGCACUAUGC GCG GAUACCUU	2819
1182	GACCACUC G CUGUACAC	1045	GUGUACAG UGAUG GCAUGCACUAUGC GCG GAGUGGUC	2820
1234	GGUAUUUU G AGGUGAUC	1046	GAUCACCU UGAUG GCAUGCACUAUGC GCG AUAAUACC	2821
1239	UAUGAGGU G AUCAUUGU	1047	ACAAUGAU UGAUG GCAUGCACUAUGC GCG ACCUCAUA	2822
1248	AUCAUUGU G CGGGUGGA	1048	UCCACCCG UGAUG GCAUGCACUAUGC GCG ACAAUGAU	2823
1275	CAGGAUCU G AAAUUGGA	1049	UCCAUUUU UGAUG GCAUGCACUAUGC GCG AGAUCCUG	2824
1286	AAUGGACU G CAAGGAGU	1050	ACUCCUUG UGAUG GCAUGCACUAUGC GCG AGUCCAUU	2825
1303	ACAACTAU G ACAAGAGC	1051	GCUCUUGU UGAUG GCAUGCACUAUGC GCG AUAGUUGU	2826
1344	CUUCGUUU G CCCAAGAA	1052	UUCUUUGG UGAUG GCAUGCACUAUGC GCG AAACGAAG	2827
1360	AAUGUUUU G AAGCUUGCA	1053	UGCAGCUU UGAUG GCAUGCACUAUGC GCG AAACACUU	2828
1366	UUGAAGCU G CAGUCAAA	1054	UUUGACUG UGAUG GCAUGCACUAUGC GCG AGCUUCAA	2829
1411	AGUUCCCU G AUGGUUUC	1055	GAACCAU UGAUG GCAUGCACUAUGC GCG AGGGAACU	2830
1442	GCUGGUGU G CUGGCAAG	1056	CUUGCCAG UGAUG GCAUGCACUAUGC GCG ACACCAGC	2831
1504	UAAUGGGU G AGGUUACC	1057	GGUAACCU UGAUG GCAUGCACUAUGC GCG ACCCAUUA	2832

Table 20

1526	GUCCUUC G CAUCACCA	1058	UGUGAUG UGAUG GCAUGCACUAUGC GCG GGAAGGAC	2833
1542	AUCCUUC G CAGCAUA	1059	UAUUGCUG UGAUG GCAUGCACUAUGC GCG GGAAGGAU	2834
1554	CAAUACCU G CGGCCAGU	1060	ACUGGCCG UGAUG GCAUGCACUAUGC GCG AGGUAUTUG	2835
1588	CCCAAGAC G ACUGUAC	1061	GUAAACAG UGAUG GCAUGCACUAUGC GCG GUCUUGGG	2836
1603	ACAAAGUU G CCAUCUCA	1062	UGAGAUGG UGAUG GCAUGCACUAUGC GCG AAACUUGU	2837
1672	UUGUCUUU G AUCGGGCC	1063	GGCCCGAU UGAUG GCAUGCACUAUGC GCG AAAGACAA	2838
1682	UCGGGCC G AAAACGAA	1064	UUCGUUUU UGAUG GCAUGCACUAUGC GCG GGGCCCCA	2839
1688	CCGAAAC G AAUUGGU	1065	AGCCAAUU UGAUG GCAUGCACUAUGC GCG GUUUUCGG	2840
1699	UUGGCUUU G CUGUCAGC	1066	GCUGACAG UGAUG GCAUGCACUAUGC GCG AAAGCCAA	2841
1708	CUGUCAGC G CUUGCCAU	1067	AUGGCAAG UGAUG GCAUGCACUAUGC GCG GCUGACAG	2842
1712	CAGGCUU G CCAUGUGC	1068	GCACAUUG UGAUG GCAUGCACUAUGC GCG AAGCGCUG	2843
1719	UGCCAUU G CACGAUGA	1069	UCAUCGUG UGAUG GCAUGCACUAUGC GCG ACAUGGCA	2844
1723	AUGUGCAC G AUGAGUUC	1070	GAACUCAU UGAUG GCAUGCACUAUGC GCG GUGCACAU	2845
1726	UGCACGAU G AGUUCAGG	1071	CCUGAACU UGAUG GCAUGCACUAUGC GCG AUCUGUCA	2846
1807	AGACAGAU G AGUCAACC	1072	GGUUGACU UGAUG GCAUGCACUAUGC GCG AUCUGUCU	2847
1821	ACCCUCAU G ACCAUAGC	1073	GCUAUGGU UGAUG GCAUGCACUAUGC GCG AUGAGGGU	2848
1843	UCAUGGCU G CCAUCUGC	1074	GCAGAUGG UGAUG GCAUGCACUAUGC GCG AGCCAUGA	2849
1850	UGCCAUCU G CGCCUCU	1075	AGAGGGCG UGAUG GCAUGCACUAUGC GCG AGAUGGCA	2850
1852	CCAUCUGC G CCUCUCUC	1076	GAAGAGGG UGAUG GCAUGCACUAUGC GCG GCAGAUGG	2851
1863	CUUCUCAU G CUGGCCAU	1077	AGUGGCAG UGAUG GCAUGCACUAUGC GCG AUGAAGAG	2852
1866	UUAUGUCU G CCACUCUG	1078	CAGAGUGG UGAUG GCAUGCACUAUGC GCG AGCAUGAA	2853
1874	GCCACUCU G CCUCAUGG	1079	CCAUGAGG UGAUG GCAUGCACUAUGC GCG AGAGUGGC	2854
1895	UCAGUGGC G CUGCCUCC	1080	GGAGGCAG UGAUG GCAUGCACUAUGC GCG GCCACUGA	2855
1898	GUGGCGCU G CCUGCCGU	1081	AGCGGAGG UGAUG GCAUGCACUAUGC GCG AGCGCCAC	2856
1904	CUGCCUCC G CUGCCUGC	1082	GCAGGCAG UGAUG GCAUGCACUAUGC GCG GGAGGCAG	2857
1907	CCUCCGCU G CCUGCGCC	1083	GGCGCAGG UGAUG GCAUGCACUAUGC GCG AGCGGAGG	2858
1911	CGCUGCCU G CGCCAGCA	1084	UGCUGGCG UGAUG GCAUGCACUAUGC GCG AGGCAGCG	2859
1913	CUGCCUGC G CCAGCAGC	1085	GCUGCUGG UGAUG GCAUGCACUAUGC GCG GCAGGCAG	2860
1924	AGCAGCAU G AUGACUUU	1086	AAAGUCAU UGAUG GCAUGCACUAUGC GCG AUGCUGCU	2861
1927	AGCAUGAU G ACUUUGCU	1087	AGCAAAGU UGAUG GCAUGCACUAUGC GCG AUCAUGCU	2862
1933	AUGACUUU G CUGAUGAC	1088	GUCAUCAG UGAUG GCAUGCACUAUGC GCG AAAGUCAU	2863
1936	ACUUUGCU G AUGACAUC	1089	GAUGUCAU UGAUG GCAUGCACUAUGC GCG AGCAAAAGU	2864
1939	UUGCUGAU G ACAUCUCC	1090	GGAGAUGU UGAUG GCAUGCACUAUGC GCG AUCAGCAA	2865
1950	AUCUCCCU G CUGAAGUG	1091	CACUUCAG UGAUG GCAUGCACUAUGC GCG AGGGAGAU	2866

Table 20

1953	UCCUGCU G AAGUGAGG	1092	CCUCACUU UGAUG GCAUGCACUAUGC GCG AGCAGGGA	2867
1958	GCUGAAGU G AGGAGGCC	1093	GGCCUCCU UGAUG GCAUGCACUAUGC GCG ACUCAGC	2868
2087	CACCAAAU G CCUCUGCC	1094	GGCAGAGG UGAUG GCAUGCACUAUGC GCG AUUUGGUG	2869
2093	AUGCCUCU G CCUUGAUG	1095	CAUCAAGG UGAUG GCAUGCACUAUGC GCG AGAGCAU	2870
2098	UCUGCCUU G AUGGAGAA	1096	UUCUCCAU UGAUG GCAUGCACUAUGC GCG AAGGAGA	2871
2179	AGCACUCU G CUGGGGG	1097	CCGGCCAG UGAUG GCAUGCACUAUGC GCG AGAGUCU	2872
2227	GAATUCU G CUGCUUGA	1098	UCAAGCAG UGAUG GCAUGCACUAUGC GCG AGAATUUC	2873
2230	AUUCUGCU G CUUGAAAC	1099	GUUUCAAG UGAUG GCAUGCACUAUGC GCG AGCAGAAU	2874
2234	UGCUGCUU G AAACUUGA	1100	UGAAGUUU UGAUG GCAUGCACUAUGC GCG AAGCAGCA	2875
2248	UCAGCCCU G AACUUUG	1101	CAAAGGUU UGAUG GCAUGCACUAUGC GCG AGGGCUGA	2876
2329	CAUCACAC G CAGGUUAC	1102	GUAAACUG UGAUG GCAUGCACUAUGC GCG GUGUGAUG	2877
2393	GUUUCUU G CUGGCCAA	1103	UUGGCCAG UGAUG GCAUGCACUAUGC GCG AGGGAAC	2878
2419	GAGAGAU G CACAGUUU	1104	AAACUGUG UGAUG GCAUGCACUAUGC GCG AUCCUCUC	2879
2428	CACAGUUU G CUUUUUG	1105	GCAAAUAG UGAUG GCAUGCACUAUGC GCG AAACUGUG	2880
2435	UGCUAUUU G CUUUAGAG	1106	CUCUAAAG UGAUG GCAUGCACUAUGC GCG AAUAAGCA	2881
2476	ACAUUGGU G CAAAGAUU	1107	AAUCUUUG UGAUG GCAUGCACUAUGC GCG ACCAAUGU	2882
2485	CAAAGAUU G CCUCUUGA	1108	UCAAGAGG UGAUG GCAUGCACUAUGC GCG AAUCUUUG	2883
2492	UGCCUCUU G AAUUA AAA	1109	UUUUAUUU UGAUG GCAUGCACUAUGC GCG AAGAGGCA	2884
219	GUGCCGAU G UAGCGGCG	1110	GCCCGCUA UGAUG GCAUGCACUAUGC GCG AUCGGCAC	2885
483	CUCCUGCU G UGGAUGGG	1111	CCCAUCCA UGAUG GCAUGCACUAUGC GCG AGCAGGAG	2886
634	GCAGCUUU G UGGAGAUG	1112	CAUCUCCA UGAUG GCAUGCACUAUGC GCG AAAGCUGC	2887
804	AGGCAGCU G UCCAGCAC	1113	GUGCUGGA UGAUG GCAUGCACUAUGC GCG AGCUGCCU	2888
835	GGAAGGGU G UGUUUGUG	1114	CACAUACA UGAUG GCAUGCACUAUGC GCG ACCUUUCC	2889
837	AAGGGUGU G UAUGUGCC	1115	GGCACAUU UGAUG GCAUGCACUAUGC GCG ACACCUU	2890
841	GUGUGUAU G UGCCCCUAC	1116	GUAGGGCA UGAUG GCAUGCACUAUGC GCG AUACACAC	2891
919	ACGUCACU G UGCGUGCC	1117	GGCACGCA UGAUG GCAUGCACUAUGC GCG AGUGACGU	2892
1100	GCAGCUUU G UGGUGCUG	1118	CAGCACCA UGAUG GCAUGCACUAUGC GCG AAAGCUGC	2893
1144	UGGCCUCU G UCGGAGGG	1119	CCUCCCGA UGAUG GCAUGCACUAUGC GCG AGAGGCCA	2894
1185	CACUCGCU G UACACAGG	1120	CCUGUGUA UGAUG GCAUGCACUAUGC GCG AGCGAGUG	2895
1246	UGAUCAUU G UGCGGGUG	1121	CACCCGCA UGAUG GCAUGCACUAUGC GCG AAUGAUA	2896
1315	AGAGCAUU G UGGACAGU	1122	ACUGUCCA UGAUG GCAUGCACUAUGC GCG AAUGCUCU	2897
1356	AAGAAAGU G UUUGAAGC	1123	GCUUCAAA UGAUG GCAUGCACUAUGC GCG ACUUUCUU	2898
1440	CAGCUGGU G UGCUGGCA	1124	UGCCAGCA UGAUG GCAUGCACUAUGC GCG ACCAGCUG	2899
1570	UGGAAGAU G UGGCCACG	1125	CGUGGCCA UGAUG GCAUGCACUAUGC GCG AUCUUCCA	2900

Table 20

1592	AGAGGACU G UUAACAAGU	1126	ACUUGUAA UGAUG GCAUGGCACUAUGC GCG AGUCGUCU	2901
1630	CGGGCACU G UUAUGGGA	1127	UCCCAUAA UGAUG GCAUGGCACUAUGC GCG AGUGCCCG	2902
1642	UGGGAGCU G UUAUCAUG	1128	CAUGAUA UGAUG GCAUGGCACUAUGC GCG AGCUCCCA	2903
1666	UCUACGUU G UCUUUGAU	1129	AUCAAGA UGAUG GCAUGGCACUAUGC GCG AACGUAGA	2904
1702	GCUUUGCU G UCAGGCGU	1130	AGCGUGA UGAUG GCAUGGCACUAUGC GCG AGCAAAGC	2905
1717	CUUGCCAU G UGCACGAU	1131	AUCGUGCA UGAUG GCAUGGCACUAUGC GCG AUGGCAAG	2906
1759	GCCUUUU G UCACCUUG	1132	CAAGGUGA UGAUG GCAUGGCACUAUGC GCG AAAAGGGC	2907
1781	GGAAGACU G UGGCUACA	1133	UGUAGCCA UGAUG GCAUGGCACUAUGC GCG AGUCUUCC	2908
1834	UAGCCUUA G UCAUGGCU	1134	AGCCAUGA UGAUG GCAUGGCACUAUGC GCG AUAGGCUA	2909
1884	CUCAUGGU G UGUCAGUG	1135	CACUGACA UGAUG GCAUGGCACUAUGC GCG ACCAUGAG	2910
1886	CAUGGUGU G UCAGUGGC	1136	GCCACUGA UGAUG GCAUGGCACUAUGC GCG ACACCAUG	2911
2048	UGGCACCU G UGGCCAGA	1137	UCUGGCCA UGAUG GCAUGGCACUAUGC GCG AGGUGCCA	2912
2139	CAGGGACU G UACCUGUA	1138	UACAGGUA UGAUG GCAUGGCACUAUGC GCG AGUCCUG	2913
2145	CUGUACCU G UAGGAAAC	1139	GUUUCUA UGAUG GCAUGGCACUAUGC GCG AGGUACAG	2914
2256	GAACCUUU G UCCACCAU	1140	AUGGUGGA UGAUG GCAUGGCACUAUGC GCG AAAGGUUC	2915
2346	CUUGGCGU G UGUCCUG	1141	CAGGGACA UGAUG GCAUGGCACUAUGC GCG ACGCCAAG	2916
2348	UGGCGUGU G UCCUGUG	1142	CACAGGGA UGAUG GCAUGGCACUAUGC GCG ACACGCCA	2917
2354	GUGUCCCU G UGUACCC	1143	GGGUACCA UGAUG GCAUGGCACUAUGC GCG AGGGACAC	2918
2385	CCAAGCUU G UUUCCUG	1144	CAGGGAUA UGAUG GCAUGGCACUAUGC GCG AAGCUUGG	2919
2453	CAGGGACU G UAUAAACA	1145	UGUUUAUA UGAUG GCAUGGCACUAUGC GCG AGUCCUG	2920

Input Sequence = AF190725. Cut Site = G/.

Stem Length = 8. Core Sequence = UGAUG GCAUGGCACUAUGC GCG

AF190725 (Homo sapiens beta-site APP cleaving enzyme (BACE) mRNA; 2526 bp)

Table 21

Table 21: Human BACE Zinzyme Ribozyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
11	ACGCGUCC G CAGCCCGC	960	GCGGGCUG GCCGAAAGGCGAGUCAAGGUCU GGACGCGU	2921
18	CGCAGCCC G CCCGGGAG	961	CUCCCGGG GCCGAAAGGCGAGUCAAGGUCU GGGCUGCG	2922
29	CGGGAGCU G CGAGCCGC	962	GCGGCUCG GCCGAAAGGCGAGUCAAGGUCU AGCUCCCG	2923
36	UGCGAGCC G CGAGCUGG	964	CCAGCUCG GCCGAAAGGCGAGUCAAGGUCU GGCUCGCA	2924
69	CAGCCAAC G CAGCCGCA	967	UGCGCUG GCCGAAAGGCGAGUCAAGGUCU GUUGGCGU	2925
75	ACGCGACC G CAGGAGCC	968	GGCUCUG GCCGAAAGGCGAGUCAAGGUCU GGCUGCGU	2926
94	GAGCCCUU G CCCUGCC	969	GGCAGGGG GCCGAAAGGCGAGUCAAGGUCU AAGGGCUC	2927
100	UUGCCCUU G CCCGCGCC	970	GGCGCGGG GCCGAAAGGCGAGUCAAGGUCU AGGGGCAA	2928
104	CCCUGCCC G CGCGCGCG	971	CGGCGGGG GCCGAAAGGCGAGUCAAGGUCU GGGCAGGG	2929
106	CUGCCCGC G CCGCGGCC	972	GGCGGGGG GCCGAAAGGCGAGUCAAGGUCU GCGGGCAG	2930
109	CCCGCGCC G CCGCCGCG	973	GCGGGGGG GCCGAAAGGCGAGUCAAGGUCU GCGCGGGG	2931
112	GCGCCGCC G CCGCGCGG	974	CCGGCGGG GCCGAAAGGCGAGUCAAGGUCU GCGCGCGC	2932
116	CGCCGCCG G CCGGGGGG	975	CCCCCGGG GCCGAAAGGCGAGUCAAGGUCU GGGCGGGG	2933
137	GGGAAGCC G CCACCGGC	976	GCCGGUGG GCCGAAAGGCGAGUCAAGGUCU GGCUCUCC	2934
148	ACCGGCCG G CCAUGCCC	977	GGGCAUGG GCCGAAAGGCGAGUCAAGGUCU GGGCCGGU	2935
153	CCCGCCAU G CCCGCCCC	978	GGGGCGGG GCCGAAAGGCGAGUCAAGGUCU AUGGCGGG	2936
157	CCAUGCCC G CCCUCCCC	979	GGGAGGGG GCCGAAAGGCGAGUCAAGGUCU GGGCAUGG	2937
172	CCAGCCCC G CCGGGAGC	980	GCUCGCGG GCCGAAAGGCGAGUCAAGGUCU GGGGCGUG	2938
183	GGGAGCCC G CGCCCGCU	981	AGCGGGCG GCCGAAAGGCGAGUCAAGGUCU GGGCUCCC	2939
185	GAGCCCGC G CCCGCGUC	982	GCAGCGGG GCCGAAAGGCGAGUCAAGGUCU GCGGGCUC	2940
189	CCGCGCCC G CUGCCAG	983	CUGGGCAG GCCGAAAGGCGAGUCAAGGUCU GGGCGCGG	2941
192	CGCCCGCU G CCCAGGCU	984	AGCCUGGG GCCGAAAGGCGAGUCAAGGUCU AGCGGGCG	2942
205	GGCUGGCC G CCGCGGUG	985	CACGGCGG GCCGAAAGGCGAGUCAAGGUCU GGCCAGCC	2943
208	UGGCCGCC G CCGUGCCG	986	CGGCACGG GCCGAAAGGCGAGUCAAGGUCU GCGGCCCA	2944
213	GCCGCGCU G CCGAUGUA	987	UACAUCGG GCCGAAAGGCGAGUCAAGGUCU ACGGCGGC	2945
250	UCUCCCUU G CUCCCGUG	989	CACGGGAG GCCGAAAGGCGAGUCAAGGUCU AGGGGAGA	2946
258	GCUCGCGU G CUCUGCGG	990	CCGAGAG GCCGAAAGGCGAGUCAAGGUCU ACGGGAGC	2947
263	CGUGUCU G CGGAUCUC	991	GAGAUCCG GCCGAAAGGCGAGUCAAGGUCU AGAGCACG	2948
280	CCCUGACC G CUCUCCAC	993	GUGGAGAG GCCGAAAGGCGAGUCAAGGUCU GGUCAGGG	2949
320	AGGGCCCU G CAGGCCCU	994	AGGGCCUG GCCGAAAGGCGAGUCAAGGUCU AGGGCCCU	2950
340	GUCUGAU G CCCCAG	996	CUUGGGGG GCCGAAAGGCGAGUCAAGGUCU AUCAGGAC	2951
397	GGGAGGC G CCAGGGAC	998	GUCGUGG GCCGAAAGGCGAGUCAAGGUCU GCGGCCC	2952
420	GGGCCAGU G CGAGCCCA	999	UGGGCUCG GCCGAAAGGCGAGUCAAGGUCU ACUGGCCC	2953
468	CAAGCCCU G CCCUGGCU	1002	AGCCAGGG GCCGAAAGGCGAGUCAAGGUCU AGGGCUUG	2954
480	UGGCUCCU G CUGUGGAU	1003	AUCCACAG GCCGAAAGGCGAGUCAAGGUCU AGGAGCCA	2955
493	GGAUGGGC G CGGGAGUG	1004	CACUCCCG GCCGAAAGGCGAGUCAAGGUCU GCCAUCC	2956
501	GCGGGAGU G CUGCCUGC	1005	GCAGGCAG GCCGAAAGGCGAGUCAAGGUCU ACUCCCGC	2957
504	GGAGUGCU G CCUGCCCA	1006	UGGGCAGG GCCGAAAGGCGAGUCAAGGUCU AGCACUCC	2958
508	UGCUGCCU G CCCACGGC	1007	GCCGUGGG GCCGAAAGGCGAGUCAAGGUCU AGGCAGCA	2959
537	AUCCGGCU G CCCUGCG	1008	CGCAGGGG GCCGAAAGGCGAGUCAAGGUCU AGCCGAU	2960
543	CUGCCCUU G CGCAGCGG	1009	CCGUGCG GCCGAAAGGCGAGUCAAGGUCU AGGGCAG	2961
545	GCCCCUGC G CAGCGGCC	1010	GGCCGCG GCCGAAAGGCGAGUCAAGGUCU GCAGGGGC	2962
562	UGGGGGG G CCCCCUG	1011	CAGGGGGG GCCGAAAGGCGAGUCAAGGUCU GCCCCCA	2963
576	CUGGGGCU G CGGUGGCC	1012	GGCAGCCG GCCGAAAGGCGAGUCAAGGUCU AGCCCCAG	2964
582	CUGCGGCU G CCCCAGGA	1013	UCCCGGGG GCCGAAAGGCGAGUCAAGGUCU AGCCGAG	2965
708	AGCCCCC G CAGACGCU	1019	AGCGUCUG GCCGAAAGGCGAGUCAAGGUCU GGGGGGCU	2966
714	CCGAGAC G CUACAACU	1020	AUGUUGAG GCCGAAAGGCGAGUCAAGGUCU GUCUGCGG	2967

Table 21

751	GUAACUUU G CAGUGGGU	1021	ACCCACUG GCCGAAAGGCGAGUCAAGGUCU AAAGUUAC	2968
760	CAGUGGGU G CUGCCCCC	1022	GGGGGCAG GCCGAAAGGCGAGUCAAGGUCU ACCCACUG	2969
763	UGGGUGCU G CCCCCAC	1023	GUGGGGGG GCCGAAAGGCGAGUCAAGGUCU AGCACCCA	2970
780	CCCUCCU G CAUCGCUA	1024	UAGCGAUG GCCGAAAGGCGAGUCAAGGUCU AGGAAGGG	2971
785	CCUGCAUC G CUACUACC	1025	GGUAGUAG GCCGAAAGGCGAGUCAAGGUCU GAUGCAGG	2972
843	GUGUAUGU G CCCUACAC	1026	GUGUAGGG GCCGAAAGGCGAGUCAAGGUCU ACAUACAC	2973
921	GUCACUGU G CGUGCCAA	1028	UUGGCACG GCCGAAAGGCGAGUCAAGGUCU ACAGUGAC	2974
925	CUGUGCGU G CCAACAUU	1029	AAUGUUGG GCCGAAAGGCGAGUCAAGGUCU ACGCACAG	2975
934	CCAACAUU G CUGCCAUC	1030	GAUGGCAG GCCGAAAGGCGAGUCAAGGUCU AAUGUUGG	2976
937	ACAUUGCU G CCAUCACU	1031	AGUGAUGG GCCGAAAGGCGAGUCAAGGUCU AGCAAUGU	2977
1006	UGGCCUUAU G CUGAGAUU	1033	AAUCUCAG GCCGAAAGGCGAGUCAAGGUCU AUAGGCCA	2978
1015	CUGAGAUU G CCAGGCCU	1035	AGGCCUGG GCCGAAAGGCGAGUCAAGGUCU AAUCUCAG	2979
1092	UUCUCCU G CAGCUUG	1039	CAAAGCUG GCCGAAAGGCGAGUCAAGGUCU AGGGAGAA	2980
1105	UUUGUGGU G CUGGCUUC	1040	GAAGCCAG GCCGAAAGGCGAGUCAAGGUCU ACCACAAA	2981
1134	UCUGAAGU G CUGGCCUC	1042	GAGGCCAG GCCGAAAGGCGAGUCAAGGUCU ACUUCAGA	2982
1182	GACCACUC G CUGUACAC	1045	GUGUACAG GCCGAAAGGCGAGUCAAGGUCU GAGUGGUC	2983
1248	AUCAUUGU G CGGGUGGA	1048	UCCACCCG GCCGAAAGGCGAGUCAAGGUCU ACAAUGAU	2984
1286	AAUGGACU G CAAGGAGU	1050	ACUCCUUG GCCGAAAGGCGAGUCAAGGUCU AGUCCAUA	2985
1344	CUUCGUUU G CCCAAGAA	1052	UUUUGGG GCCGAAAGGCGAGUCAAGGUCU AAACGAAG	2986
1366	UUGAAGU G CAGUCAA	1054	UUUGACUG GCCGAAAGGCGAGUCAAGGUCU AGCUUCAA	2987
1442	GCUGGUGU G CUGGCAAG	1056	CUUGCCAG GCCGAAAGGCGAGUCAAGGUCU ACACCAGC	2988
1526	GUCCUUC G CAUCACCA	1058	UGGUGAUG GCCGAAAGGCGAGUCAAGGUCU GGAAGGAC	2989
1542	AUCCUUC G CAGCAUA	1059	UAUUGCUG GCCGAAAGGCGAGUCAAGGUCU GGAAGGAU	2990
1554	CAAUACCU G CGGCCAGU	1060	ACUGGCCG GCCGAAAGGCGAGUCAAGGUCU AGGUUAUG	2991
1603	ACAAGUUU G CCAUCUA	1062	UGAGAUGG GCCGAAAGGCGAGUCAAGGUCU AAACUUGU	2992
1699	UUGGCUUU G CUGUCAGC	1066	GCUGACAG GCCGAAAGGCGAGUCAAGGUCU AAAGCCAA	2993
1708	CUGUCAGC G CUUGCCAU	1067	AUGGCAAG GCCGAAAGGCGAGUCAAGGUCU GCUGACAG	2994
1712	CAGCGCUU G CCAUGUGC	1068	GCACAUGG GCCGAAAGGCGAGUCAAGGUCU AAGCGCUG	2995
1719	UGCCAUGU G CACGAUGA	1069	UCAUCGUG GCCGAAAGGCGAGUCAAGGUCU ACAUGGCA	2996
1843	UCAUGGCU G CCAUCUGC	1074	GCAGAUGG GCCGAAAGGCGAGUCAAGGUCU AGCCAUGA	2997
1850	UGCCAUCU G CGCCUCU	1075	AGAGGGCG GCCGAAAGGCGAGUCAAGGUCU AGAUGGCA	2998
1852	CCAUCUGC G CCCUCUUC	1076	GAAGAGGG GCCGAAAGGCGAGUCAAGGUCU GCAGAUGG	2999
1863	CUCUUCU G CUGCCACU	1077	AGUGGCAG GCCGAAAGGCGAGUCAAGGUCU AUGAAGAG	3000
1866	UUCAUGCU G CCACUCUG	1078	CAGAGUGG GCCGAAAGGCGAGUCAAGGUCU AGCAUGAA	3001
1874	GCCACUCU G CCUCAUGG	1079	CCAUGAGG GCCGAAAGGCGAGUCAAGGUCU AGAGUGGC	3002
1895	UCAGUGGC G CUGCCUCC	1080	GGAGGCAG GCCGAAAGGCGAGUCAAGGUCU GCCACUGA	3003
1898	GUGGCGCU G CCUCCGCU	1081	AGCGGAGG GCCGAAAGGCGAGUCAAGGUCU AGCGCCAC	3004
1904	CUGCCUCC G CUGCCUGC	1082	GCAGGCAG GCCGAAAGGCGAGUCAAGGUCU GGAGGCAG	3005
1907	CCUCCGCU G CCUGCGCC	1083	GGCGCAGG GCCGAAAGGCGAGUCAAGGUCU AGCGGAGG	3006
1911	CGCUGCCU G CGCCAGCA	1084	UGCUGGCG GCCGAAAGGCGAGUCAAGGUCU AGGCAGCG	3007
1913	CUGCCUGC G CCAGCAGC	1085	GCUGCUGG GCCGAAAGGCGAGUCAAGGUCU GCAGGCAG	3008
1933	AUGACUUU G CUGAUGAC	1088	GUCAUCAG GCCGAAAGGCGAGUCAAGGUCU AAAGUCAU	3009
1950	AUCUCCU G CUGAAGUG	1091	CACUUCAG GCCGAAAGGCGAGUCAAGGUCU AGGGAGAU	3010
2087	CACCAAU G CCUCUGCC	1094	GGCAGAGG GCCGAAAGGCGAGUCAAGGUCU AUUUGGUG	3011
2093	AUGCCUCU G CCUUGAUG	1095	CAUCAAGG GCCGAAAGGCGAGUCAAGGUCU AGAGGCAU	3012
2179	AGCACUCU G CUGGCGGG	1097	CCCGCCAG GCCGAAAGGCGAGUCAAGGUCU AGAGUGCU	3013
2227	GAAAUUCU G CUGCUUGA	1098	UCAAGCAG GCCGAAAGGCGAGUCAAGGUCU AGAAUUUC	3014
2230	AUUCUGCU G CUUGAAAC	1099	GUUUCAAG GCCGAAAGGCGAGUCAAGGUCU AGCAGAAU	3015
2329	CAUCACAC G CAGGUUAC	1102	GUAACCUG GCCGAAAGGCGAGUCAAGGUCU GUGUGAUG	3016
2393	GUUUCCU G CUGGCCAA	1103	UUGGCCAG GCCGAAAGGCGAGUCAAGGUCU AGGGAAAC	3017
2419	GAGAGGAU G CACAGUUU	1104	AAACUGUG GCCGAAAGGCGAGUCAAGGUCU AUCCUCUC	3018

Table 21

2428	CACAGUUU G CUAUUUGC	1105	GCAAUAG GCCGAAAGGCGAGUCAAGGUCU AAACUGUG	3019
2435	UGCUAUUU G CUUUAGAG	1106	CUCUAAAG GCCGAAAGGCGAGUCAAGGUCU AAUAGCA	3020
2476	ACAUUGGU G CAAAGAUU	1107	AAUCUUUG GCCGAAAGGCGAGUCAAGGUCU ACCAAUGU	3021
2485	CAAAGAUU G CCUCUUGA	1108	UCAAGAGG GCCGAAAGGCGAGUCAAGGUCU AAUCUUUG	3022
219	GUGCCGAU G UAGCGGGC	1110	GCCCCUA GCCGAAAGGCGAGUCAAGGUCU AUCGGCAC	3023
483	CUCCUGCU G UGGAUUGG	1111	CCCAUCCA GCCGAAAGGCGAGUCAAGGUCU AGCAGGAG	3024
634	GCAGCUUU G UGGAGAUG	1112	CAUCUCCA GCCGAAAGGCGAGUCAAGGUCU AAAGCUGC	3025
804	AGGCAGCU G UCCAGCAC	1113	GUGCUGGA GCCGAAAGGCGAGUCAAGGUCU AGCUGCCU	3026
835	GGAAGGGU G UGU AUGUG	1114	CACAUACA GCCGAAAGGCGAGUCAAGGUCU ACCCUUCC	3027
837	AAGGGUGU G UAUGUGCC	1115	GGCACAUU GCCGAAAGGCGAGUCAAGGUCU ACACCCUU	3028
841	GUGUGUAU G UGCCCUC	1116	GUAGGGCA GCCGAAAGGCGAGUCAAGGUCU AUACACAC	3029
919	ACGUCACU G UGCGUGCC	1117	GGCACGCA GCCGAAAGGCGAGUCAAGGUCU AGUGACGU	3030
1100	GCAGCUUU G UGGUGCUG	1118	CAGCACCA GCCGAAAGGCGAGUCAAGGUCU AAAGCUGC	3031
1144	UGGCCUCU G UCGGAGGG	1119	CCCUCGGA GCCGAAAGGCGAGUCAAGGUCU AGAGGCCA	3032
1185	CACUCGCU G UACACAGG	1120	CCUGUGUA GCCGAAAGGCGAGUCAAGGUCU AGCGAGUG	3033
1246	UGAUCAUU G UGCGGGUG	1121	CACCCGCA GCCGAAAGGCGAGUCAAGGUCU AAUGAUA	3034
1315	AGAGCAUU G UGGACAGU	1122	ACUGUCCA GCCGAAAGGCGAGUCAAGGUCU AAUGUCU	3035
1356	AAGAAAGU G UUGAAGC	1123	GCUCAAA GCCGAAAGGCGAGUCAAGGUCU ACUUUCUU	3036
1440	CAGCUGGU G UGCGGCA	1124	UGCGAGCA GCCGAAAGGCGAGUCAAGGUCU ACCAGCUG	3037
1570	UGGAAGAU G UGGCCACG	1125	CGUGGCCA GCCGAAAGGCGAGUCAAGGUCU AUCUCCA	3038
1592	AGACGACU G UUAACAAGU	1126	ACUUGUAA GCCGAAAGGCGAGUCAAGGUCU AGUCGUCU	3039
1630	CGGGCACU G UUAUGGGA	1127	UCCCAUAA GCCGAAAGGCGAGUCAAGGUCU AGUGCCCG	3040
1642	UGGGAGCU G UUAUCAUG	1128	CAUGAUAA GCCGAAAGGCGAGUCAAGGUCU AGCUCCCA	3041
1666	UCUACGUU G UCUUUGAU	1129	AUCAAGA GCCGAAAGGCGAGUCAAGGUCU AACGUAGA	3042
1702	GUUUGCU G UCAGCGCU	1130	AGCGCUGA GCCGAAAGGCGAGUCAAGGUCU AGCAAAGC	3043
1717	CUUGCCAU G UGCACGAU	1131	AUCGUGCA GCCGAAAGGCGAGUCAAGGUCU AUGGCAAG	3044
1759	GCCUUUUU G UCACCUUG	1132	CAAGGUGA GCCGAAAGGCGAGUCAAGGUCU AAAAGGGC	3045
1781	GGAAGACU G UGGCUACA	1133	UGUAGCCA GCCGAAAGGCGAGUCAAGGUCU AGUCUCC	3046
1834	UAGCCUAU G UCAUGGCU	1134	AGCCAUGA GCCGAAAGGCGAGUCAAGGUCU AUAGGCUA	3047
1884	CUCAUGGU G UGUCAGUG	1135	CACUGACA GCCGAAAGGCGAGUCAAGGUCU ACCAUGAG	3048
1886	CAUGGUGU G UCAGUGGC	1136	GCCACUGA GCCGAAAGGCGAGUCAAGGUCU ACACCAUG	3049
2048	UGGACCU G UGGCCAGA	1137	UCUGGCCA GCCGAAAGGCGAGUCAAGGUCU AGGUGCCA	3050
2139	CAGGGACU G UACCUGUA	1138	UACAGGUA GCCGAAAGGCGAGUCAAGGUCU AGUCCUG	3051
2145	CUGUACCU G UAGGAAAC	1139	GUUUCUA GCCGAAAGGCGAGUCAAGGUCU AGGUACAG	3052
2256	GAACCUUU G UCCACCAU	1140	AUGGUGGA GCCGAAAGGCGAGUCAAGGUCU AAAGGUUC	3053
2346	CUUGGCGU G UGUCCUG	1141	CAGGGACA GCCGAAAGGCGAGUCAAGGUCU ACGCCAAG	3054
2348	UGGCGUGU G UCCCUGUG	1142	CACAGGGA GCCGAAAGGCGAGUCAAGGUCU ACACGCCA	3055
2354	GUGUCCCU G UGGUACCC	1143	GGGUACCA GCCGAAAGGCGAGUCAAGGUCU AGGGACAC	3056
2385	CCAAGCUU G UUUCCUG	1144	CAGGGAAA GCCGAAAGGCGAGUCAAGGUCU AAGCUUGG	3057
2453	CAGGGACU G UAUAAACA	1145	UGUUUAUA GCCGAAAGGCGAGUCAAGGUCU AGUCCUG	3058
14	CGUCCGA G CCCGCCG	1146	CGGGCGGG GCCGAAAGGCGAGUCAAGGUCU UGCGGACG	3059
26	GCCCGGA G CUGCGAGC	1147	GCUCGCG GCCGAAAGGCGAGUCAAGGUCU UCCCGGGC	3060
33	AGCUGCGA G CCGCGAGC	1148	GCUCGCG GCCGAAAGGCGAGUCAAGGUCU UCGCAGCU	3061
40	AGCCGCGA G CUGGAUUA	1149	UAAUCCAG GCCGAAAGGCGAGUCAAGGUCU UCGCGGCU	3062
51	GGAUUAUG G UGGCCUGA	1150	UCAGGCCA GCCGAAAGGCGAGUCAAGGUCU CAUAAUCC	3063
54	UUAUGGUG G CCUGAGCA	1151	UGCUCAG GCCGAAAGGCGAGUCAAGGUCU CACCAUAA	3064
60	UGGCCUGA G CAGCCAAC	1152	GUUGGUG GCCGAAAGGCGAGUCAAGGUCU UGAGGCCA	3065
63	CCUGAGCA G CCAACGCA	1153	GUUGGUG GCCGAAAGGCGAGUCAAGGUCU UGAGGCCA	3066
72	CCAACGCA G CCGCAGGA	1154	UCCUGCGG GCCGAAAGGCGAGUCAAGGUCU UGCGUUGG	3067
81	CCGCAGGA G CCGGAGC	1155	GCUCGCG GCCGAAAGGCGAGUCAAGGUCU UCCUGCGG	3068
88	AGCCCGGA G CCCUUGCC	1156	GGCAAGGG GCCGAAAGGCGAGUCAAGGUCU UCCGGGCU	3069

Table 21

134	CCAGGGAA G CCGCCACC	1157	GGUGGCGG GCCGAAAGGCGAGUCAAGGUCU	UUCCCUUGG	3070
144	CGCCACCG G CCCGCCAU	1158	AUGGCGGG GCCGAAAGGCGAGUCAAGGUCU	CGGUGGCG	3071
167	CCCUCCCA G CCCCGCCG	1159	CGGCGGGG GCCGAAAGGCGAGUCAAGGUCU	UGGGAGGG	3072
179	CGCCGGGA G CCCCGGCC	1160	GGCGCGGG GCCGAAAGGCGAGUCAAGGUCU	UCCCGGCG	3073
198	CUGCCAG G CUGGCCGC	1161	GCGGCCAG GCCGAAAGGCGAGUCAAGGUCU	CUGGGCAG	3074
202	CCAGGCUG G CCGCCGCC	1162	GGCGCGGG GCCGAAAGGCGAGUCAAGGUCU	CAGCCUGG	3075
211	CCGCCGCC G UGCCGAUG	1163	CAUCGGCA GCCGAAAGGCGAGUCAAGGUCU	GGCGGCGG	3076
222	CCGAUGUA G CGGGCUCC	1164	GGAGCCCG GCCGAAAGGCGAGUCAAGGUCU	UACAUCGG	3077
226	UGUAGCGG G CUCCGGAU	1165	AUCCGGAG GCCGAAAGGCGAGUCAAGGUCU	CCGCUACA	3078
239	GGAUCCCA G CCUCUCCC	1166	GGGAGAGG GCCGAAAGGCGAGUCAAGGUCU	UGGGAUCC	3079
256	CUGCUCC G UGCUCUGC	1167	GCAGAGCA GCCGAAAGGCGAGUCAAGGUCU	GGGAGCAG	3080
290	UCUCCACA G CCCGACC	1168	GGUCCGGG GCCGAAAGGCGAGUCAAGGUCU	UGUGGAGA	3081
304	ACCCGGGG G CUGGCCCA	1169	UGGGCCAG GCCGAAAGGCGAGUCAAGGUCU	CCCCGGGU	3082
308	GGGGGUG G CCAGGGC	1170	GCCUGGG GCCGAAAGGCGAGUCAAGGUCU	CAGCCCCC	3083
315	GGCCAGG G CCUGCAG	1171	CUGCAGGG GCCGAAAGGCGAGUCAAGGUCU	CCUGGGCC	3084
324	CCUGCAG G CCUGGCG	1172	CGCCAGGG GCCGAAAGGCGAGUCAAGGUCU	CUGCAGGG	3085
330	AGGCCUG G CGUCCUGA	1173	UCAGGACG GCCGAAAGGCGAGUCAAGGUCU	CAGGGCCU	3086
332	GCCUGGC G UCCUGAUG	1174	CAUCAGGA GCCGAAAGGCGAGUCAAGGUCU	GCCAGGGC	3087
348	GCCCCAA G CUCCUCU	1175	AGAGGGAG GCCGAAAGGCGAGUCAAGGUCU	UUGGGGGC	3088
365	CCUGAGAA G CCACCAGC	1176	GCUGGUGG GCCGAAAGGCGAGUCAAGGUCU	UUCUCAGG	3089
372	AGCCACCA G CACCACC	1177	GGGUGGUG GCCGAAAGGCGAGUCAAGGUCU	UGGUGGCU	3090
391	ACUUGGG G CAGGCGCC	1178	GGCGCCUG GCCGAAAGGCGAGUCAAGGUCU	CCCCAAGU	3091
395	GGGGGAG G CGCCAGGG	1179	CCCUGGCG GCCGAAAGGCGAGUCAAGGUCU	CUGCCCCC	3092
410	GGACGGAC G UGGCCAG	1180	CUGGCCCA GCCGAAAGGCGAGUCAAGGUCU	GUCCGUCC	3093
414	GGACGUGG G CCAGUGCG	1181	CGCACUGG GCCGAAAGGCGAGUCAAGGUCU	CCACGUCC	3094
418	GUGGGCCA G UGCGAGCC	1182	GGCUCGCA GCCGAAAGGCGAGUCAAGGUCU	UGGCCCAC	3095
424	CAGUGCGA G CCAGAGG	1183	CCUCUGGG GCCGAAAGGCGAGUCAAGGUCU	UCGCACUG	3096
433	CCCAGAGG G CCGAAGG	1184	CCUUCGGG GCCGAAAGGCGAGUCAAGGUCU	CCUCUGGG	3097
441	GCCCGAAG G CCGGGGCC	1185	GGCCCCGG GCCGAAAGGCGAGUCAAGGUCU	CUUCGGGC	3098
447	AGGCCGGG G CCCACCAU	1186	AUGGUGGG GCCGAAAGGCGAGUCAAGGUCU	CCCGGCCU	3099
457	CCACCAUG G CCCAAGCC	1187	GGCUUGGG GCCGAAAGGCGAGUCAAGGUCU	CAUGGUGG	3100
463	UGGCCAA G CCUGCCC	1188	GGGCAGGG GCCGAAAGGCGAGUCAAGGUCU	UUGGGCCA	3101
474	CUGCCUG G CUCCUGCU	1189	AGCAGGAG GCCGAAAGGCGAGUCAAGGUCU	CAGGGCAG	3102
491	GUGGAUGG G CGCGGGAG	1190	CUCCCGCG GCCGAAAGGCGAGUCAAGGUCU	CCAUCCAC	3103
499	GCGCGGGA G UGUGCCU	1191	AGGCAGCA GCCGAAAGGCGAGUCAAGGUCU	UCCCGCGC	3104
515	UGCCACG G CACCCAGC	1192	GCUGGGUG GCCGAAAGGCGAGUCAAGGUCU	CGUGGGCA	3105
522	GGACCCA G CACGGCAU	1193	AUGCCGUG GCCGAAAGGCGAGUCAAGGUCU	UGGGUGCC	3106
527	CCAGCAG G CAUCCGGC	1194	GCCGGAUG GCCGAAAGGCGAGUCAAGGUCU	CGUGCUGG	3107
534	GGCAUCCG G CUGCCCU	1195	AGGGGCG GCCGAAAGGCGAGUCAAGGUCU	CGGAUGCC	3108
548	CCUGCGCA G CGGCCUGG	1196	CCAGCCG GCCGAAAGGCGAGUCAAGGUCU	UGCGCAGG	3109
551	GCGCAGCG G CCUGGGG	1197	CCCCCAGG GCCGAAAGGCGAGUCAAGGUCU	CGCCCGCG	3110
560	CCUGGGGG G CGCCCCC	1198	GGGGGGCG GCCGAAAGGCGAGUCAAGGUCU	CCCCCAGG	3111
573	CCCCUGGG G CUGCGGCU	1199	AGCCGCAG GCCGAAAGGCGAGUCAAGGUCU	CCCAGGGG	3112
579	GGGUGCG G CUGCCCCG	1200	CGGGGCAG GCCGAAAGGCGAGUCAAGGUCU	CGCAGCCC	3113
603	GACGAAGA G CCCGAGGA	1201	UCCUCGGG GCCGAAAGGCGAGUCAAGGUCU	UCUUCGUC	3114
612	CCCGAGGA G CCCGCCG	1202	CGGCCGGG GCCGAAAGGCGAGUCAAGGUCU	UCCUCGGG	3115
617	GGAGCCCG G CCGGAGGG	1203	CCUCCGG GCCGAAAGGCGAGUCAAGGUCU	CGGGCUCC	3116
626	CCGAGGG G CAGCUUG	1204	CAAAGCUG GCCGAAAGGCGAGUCAAGGUCU	CCUCCGG	3117
629	GAGGGGCA G CUUUGUGG	1205	CCACAAAG GCCGAAAGGCGAGUCAAGGUCU	UGCCCCUC	3118
643	UGGAGAUG G UGGACAAC	1206	GUUGUCCA GCCGAAAGGCGAGUCAAGGUCU	CAUCUCCA	3119
659	CCUGAGGG G CAAGUCGG	1207	CCGACUUG GCCGAAAGGCGAGUCAAGGUCU	CCUCCAGG	3120

Table 21

663	AGGGGCAA G UCGGGGCA	1208	UGCCCCGA GCCGAAAGGCGAGUCAAGGUCU UUGCCCCU	3121
669	AAGUCGGG G CAGGGCUA	1209	UAGCCUG GCCGAAAGGCGAGUCAAGGUCU CCCGACUU	3122
674	GGGGCAGG G CUACUACG	1210	CGUAGUAG GCCGAAAGGCGAGUCAAGGUCU CCUGCCCC	3123
682	GCUACUAC G UGGAGAUG	1211	CAUCUCCA GCCGAAAGGCGAGUCAAGGUCU GUAGUAGC	3124
694	AGAUGACC G UGGGCAGC	1212	GCUGCCCA GCCGAAAGGCGAGUCAAGGUCU GGUCAUCU	3125
698	GACCGUGG G CAGCCCCC	1213	GGGGGUG GCCGAAAGGCGAGUCAAGGUCU CCACGGUC	3126
701	CGUGGGCA G CCCCCCGC	1214	GCGGGGG GCCGAAAGGCGAGUCAAGGUCU UGCCCACG	3127
727	ACAUCUG G UGGAUACA	1215	UGUAUCCA GCCGAAAGGCGAGUCAAGGUCU CAGGAUGU	3128
737	GAUACAG G CAGCAGUA	1216	UACUGCUG GCCGAAAGGCGAGUCAAGGUCU CUGUAUCC	3129
740	UACAGGCA G CAGUAACU	1217	AGUUAUCG GCCGAAAGGCGAGUCAAGGUCU UGCCUGUA	3130
743	AGGCAGCA G UAACUUUG	1218	CAAAGUUA GCCGAAAGGCGAGUCAAGGUCU UGCUGCCU	3131
754	ACUUUGCA G UGGGUGCU	1219	AGCACCCA GCCGAAAGGCGAGUCAAGGUCU UGCAAAGU	3132
758	UGCAGUGG G UGCUGCCC	1220	GGGCAGCA GCCGAAAGGCGAGUCAAGGUCU CCACUGCA	3133
798	UACCAGAG G CAGCUGUC	1221	GACAGCUG GCCGAAAGGCGAGUCAAGGUCU CUCUGGUA	3134
801	CAGAGGCA G CUGUCCAG	1222	CUGGACAG GCCGAAAGGCGAGUCAAGGUCU UGCCUCUG	3135
809	GCUGUCCA G CACAUACC	1223	GGUAUGUG GCCGAAAGGCGAGUCAAGGUCU UGGACAGC	3136
833	CCGGAAGG G UGUGUAUG	1224	CAUACACA GCCGAAAGGCGAGUCAAGGUCU CCUUCGG	3137
857	CACCCAGG G CAAGUGGG	1225	CCCACUUG GCCGAAAGGCGAGUCAAGGUCU CCUGGGUG	3138
861	CAGGGCAA G UGGGAAGG	1226	CCUUCCCA GCCGAAAGGCGAGUCAAGGUCU UUGCCUG	3139
873	GAAGGGA G CUGGGCAC	1227	GUGCCAG GCCGAAAGGCGAGUCAAGGUCU UCCCUUC	3140
878	GGAGCUGG G CACCGACC	1228	GGUCGGUG GCCGAAAGGCGAGUCAAGGUCU CCAGCUCC	3141
889	CCGACCUG G UAAGCAUC	1229	GAUGCUUA GCCGAAAGGCGAGUCAAGGUCU CAGGUCGG	3142
893	CCUGGUAA G CAUCCCCC	1230	GGGGGAUG GCCGAAAGGCGAGUCAAGGUCU UUACCAGG	3143
905	CCCCAUG G CCCCAACG	1231	CGUUGGG GCCGAAAGGCGAGUCAAGGUCU CAUGGGGG	3144
913	GCCCAAC G UCACUGUG	1232	CACAGUGA GCCGAAAGGCGAGUCAAGGUCU GUUGGGGC	3145
923	CACUGUGC G UGCCAACA	1233	UGUUGGCA GCCGAAAGGCGAGUCAAGGUCU GCACAGUG	3146
957	UCAGACAA G UUCUUAU	1234	AUGAAGAA GCCGAAAGGCGAGUCAAGGUCU UUGUCUGA	3147
971	CAUCAACG G CUCCAACU	1235	AGUUGGAG GCCGAAAGGCGAGUCAAGGUCU CGUUGAUG	3148
986	CUGGGAAG G CAUCCUGG	1236	CCAGGAUG GCCGAAAGGCGAGUCAAGGUCU CUUCCAG	3149
996	AUCCUGGG G CUGGCCUA	1237	UAGGCCAG GCCGAAAGGCGAGUCAAGGUCU CCCAGGAU	3150
1000	UGGGGUG G CCUAUGCU	1238	AGCAUAGG GCCGAAAGGCGAGUCAAGGUCU CAGCCCCA	3151
1020	AUUGCCAG G CCUGACGA	1239	UCGUCAGG GCCGAAAGGCGAGUCAAGGUCU CUGGCAAU	3152
1038	UCCUGGA G CCUUUCUU	1240	AAGAAAGG GCCGAAAGGCGAGUCAAGGUCU UCCAGGGA	3153
1057	ACUCUCUG G UAAAGCAG	1241	CUGCUUUA GCCGAAAGGCGAGUCAAGGUCU CAGAGAGU	3154
1062	CUGGUAAA G CAGACCCA	1242	UGGUCUG GCCGAAAGGCGAGUCAAGGUCU UUUACCAG	3155
1072	AGACCCAC G UUCCCAAC	1243	GUUGGGAA GCCGAAAGGCGAGUCAAGGUCU GUGGGUCU	3156
1095	UCCUGCA G CUUUGUGG	1244	CCACAAAG GCCGAAAGGCGAGUCAAGGUCU UGCAGGGA	3157
1103	GCUUUGUG G UGCUGGCU	1245	AGCCAGCA GCCGAAAGGCGAGUCAAGGUCU CACAAAGC	3158
1109	UGGUGCUG G CUUCCCCC	1246	GGGGGAAG GCCGAAAGGCGAGUCAAGGUCU CAGCACCA	3159
1125	CUCAACCA G UCUGAAGU	1247	ACUUCAGA GCCGAAAGGCGAGUCAAGGUCU UGGUUGAG	3160
1132	AGUCUGAA G UGCUGGCC	1248	GGCCAGCA GCCGAAAGGCGAGUCAAGGUCU UUCAGACU	3161
1138	AAGUCUG G CCUCUGUC	1249	GACAGAGG GCCGAAAGGCGAGUCAAGGUCU CAGCACUU	3162
1154	CGGAGGGA G CAUGAUCA	1250	UGAUCAUG GCCGAAAGGCGAGUCAAGGUCU UCCUCCG	3163
1169	CAUUGGAG G UAUCGACC	1251	GGUCGAUA GCCGAAAGGCGAGUCAAGGUCU CUCCA AUG	3164
1193	GUACACAG G CAGUCUCU	1252	AGAGACUG GCCGAAAGGCGAGUCAAGGUCU CUGUGUAC	3165
1196	CACAGGCA G UCUCUGGU	1253	ACCAGAGA GCCGAAAGGCGAGUCAAGGUCU UGCCUGUG	3166
1203	AGUCUCUG G UAUACACC	1254	GGUGUAUA GCCGAAAGGCGAGUCAAGGUCU CAGAGACU	3167
1218	CCCAUCCG G CGGGAGUG	1255	CACUCCG GCCGAAAGGCGAGUCAAGGUCU CGGAUGGG	3168
1224	CGGCGGGA G UGUUAUUA	1256	UAUAUACA GCCGAAAGGCGAGUCAAGGUCU UCCGCGG	3169
1227	CGGGAGUG G UAUUAUGA	1257	UCAUAUA GCCGAAAGGCGAGUCAAGGUCU CACUCCG	3170
1237	AUAUGAG G UGAUCAUU	1258	AAUGAUCA GCCGAAAGGCGAGUCAAGGUCU CUCAUAU	3171

Table 21

1252	UUGUGCGG G UGGAGAUC	1259	GAUCUCCA GCCGAAAGGCGAGUCAAGGUCU CCGCACAA	3172
1293	UGCAAGGA G UACAACUA	1260	UAGUUGUA GCCGAAAGGCGAGUCAAGGUCU UCCUUGCA	3173
1310	UGACAAGA G CAUUGUGG	1261	CCACAAUG GCCGAAAGGCGAGUCAAGGUCU UCUUGUCA	3174
1322	UGUGGACA G UGGACCA	1262	UGGUGCCA GCCGAAAGGCGAGUCAAGGUCU UGUCCACA	3175
1325	GGACAGUG G CACCACCA	1263	UGGUGGUG GCCGAAAGGCGAGUCAAGGUCU CACUGUCC	3176
1340	CAACCUUC G UUUGCCCA	1264	UGGGCAAA GCCGAAAGGCGAGUCAAGGUCU GAAGGUUG	3177
1354	CCAAGAAA G UGUUUGAA	1265	UUCAAACA GCCGAAAGGCGAGUCAAGGUCU UUUCUUGG	3178
1363	UGUUUGAA G CUGCAGUC	1266	GACUCGAG GCCGAAAGGCGAGUCAAGGUCU UJCAAACA	3179
1369	AAGCUGCA G UCAAAUCC	1267	GGAUUUGA GCCGAAAGGCGAGUCAAGGUCU UGCAGCUU	3180
1384	CCAUCAAG G CAGCCUCC	1268	GGAGGCGU GCCGAAAGGCGAGUCAAGGUCU CUUGAUGG	3181
1387	UCAAGGCA G CCUCCUCC	1269	GGAGGAGG GCCGAAAGGCGAGUCAAGGUCU UGCCUUGA	3182
1404	ACGGAGAA G UUCCUGA	1270	UCAGGGAA GCCGAAAGGCGAGUCAAGGUCU UUCUCCGU	3183
1415	CCCUGAUG G UUUCUGG	1271	GCCAGAAA GCCGAAAGGCGAGUCAAGGUCU CAUCAGGG	3184
1422	GGUUUCUG G CUAGGAGA	1272	UCUCCUAG GCCGAAAGGCGAGUCAAGGUCU CAGAAACC	3185
1431	CUAGGAGA G CAGCUGGU	1273	ACCAGCUG GCCGAAAGGCGAGUCAAGGUCU UCUCUAG	3186
1434	GGAGAGCA G CUGGUGUG	1274	CACACCAG GCCGAAAGGCGAGUCAAGGUCU UGCUCUCC	3187
1438	AGCAGCUG G UGUGCUGG	1275	CCAGCACA GCCGAAAGGCGAGUCAAGGUCU CAGCUGCU	3188
1446	GUGUGCUG G CAAGCAGG	1276	CCUGCUUG GCCGAAAGGCGAGUCAAGGUCU CAGCACAC	3189
1450	GCUGGCAA G CAGGCACC	1277	GGUGCCUG GCCGAAAGGCGAGUCAAGGUCU UUGCCAGC	3190
1454	GCAAGCAG G CACCACCC	1278	GGGUGGUG GCCGAAAGGCGAGUCAAGGUCU CUGCUUGC	3191
1480	UUUCCCCA G UCAUCUCA	1279	UGAGAUGA GCCGAAAGGCGAGUCAAGGUCU UGGGAAAA	3192
1502	CCUAAUGG G UGAGGUUA	1280	UAACCUCA GCCGAAAGGCGAGUCAAGGUCU CCAUUAGG	3193
1507	UGGGUGAG G UUACCAAC	1281	GUUGGUAA GCCGAAAGGCGAGUCAAGGUCU CUCACCCA	3194
1518	ACCAACCA G UCCUCCG	1282	CGGAAGGA GCCGAAAGGCGAGUCAAGGUCU UGGUUGGU	3195
1545	CUUCCGCA G CAUACCU	1283	AGGUUUG GCCGAAAGGCGAGUCAAGGUCU UGCGBAAG	3196
1557	UACCUGCG G CCAGUGGA	1284	UCCACUGG GCCGAAAGGCGAGUCAAGGUCU CGCAGGUA	3197
1561	UGCGGCCA G UGGAAGAU	1285	AUCUJCCA GCCGAAAGGCGAGUCAAGGUCU UGGCCGCA	3198
1573	AAGAUGUG G CCACGUCC	1286	GGACGUGG GCCGAAAGGCGAGUCAAGGUCU CACAUCUU	3199
1578	GUGGCCAC G UCCCAAGA	1287	UCUUGGGA GCCGAAAGGCGAGUCAAGGUCU GUGGCCAC	3200
1599	UGUACAA G UUUGCAU	1288	AUGGCAAA GCCGAAAGGCGAGUCAAGGUCU UUGUAACA	3201
1614	AUCUCACA G UCAUCCAC	1289	GUGGAUGA GCCGAAAGGCGAGUCAAGGUCU UGUGAGAU	3202
1625	AUCCACGG G CACUGUUA	1290	UAACAGUG GCCGAAAGGCGAGUCAAGGUCU CCGUGGAU	3203
1639	UUUUGGGA G CUGUUAUC	1291	GAUAACAG GCCGAAAGGCGAGUCAAGGUCU UCCAUAAA	3204
1655	CAUGGAGG G CUUCUACG	1292	CGUAGAAG GCCGAAAGGCGAGUCAAGGUCU CCUCAUG	3205
1663	GCUCUAC G UUGUCUUU	1293	AAAGACAA GCCGAAAGGCGAGUCAAGGUCU GUAGAAGC	3206
1678	UUGAUCGG G CCCGAAA	1294	UUUUCGGG GCCGAAAGGCGAGUCAAGGUCU CCGAUCAA	3207
1694	ACGAAUUG G CUUUGCUG	1295	CAGCAAAG GCCGAAAGGCGAGUCAAGGUCU CAUUCUGU	3208
1706	UGCUGUCA G CGCUUGCC	1296	GGCAAGCG GCCGAAAGGCGAGUCAAGGUCU UGACAGCA	3209
1728	CACGAUGA G UUCAGGAC	1297	GUCCUGAA GCCGAAAGGCGAGUCAAGGUCU UCAUCGUG	3210
1738	UCAGGACG G CAGCGGUG	1298	CACCGCUG GCCGAAAGGCGAGUCAAGGUCU CGUCCUGA	3211
1741	GGACGCA G CGGUGGAA	1299	UUCCACCG GCCGAAAGGCGAGUCAAGGUCU UGCCGUCC	3212
1744	CGGCAGCG G UGGAAGGC	1300	GCCUJCCA GCCGAAAGGCGAGUCAAGGUCU CGCUGCCG	3213
1751	GGUGGAAG G CCCUUUG	1301	CAAAAGGG GCCGAAAGGCGAGUCAAGGUCU CUUCCACC	3214
1784	AGACUGUG G CUACAACA	1302	UGUUGUAG GCCGAAAGGCGAGUCAAGGUCU CACAGUCU	3215
1809	ACAGAUGA G UCAACCCU	1303	AGGGUUGA GCCGAAAGGCGAGUCAAGGUCU UCAUCUGU	3216
1828	UGACCAUA G CCUAUGUC	1304	GACAUAGG GCCGAAAGGCGAGUCAAGGUCU UAUGGUA	3217
1840	AUGUCAUG G CUGCCAUC	1305	GAUGGCAG GCCGAAAGGCGAGUCAAGGUCU CAUGACAU	3218
1882	GCCUCAUG G UGUGUCAG	1306	CUGACACA GCCGAAAGGCGAGUCAAGGUCU CAUGAGGC	3219
1890	GUGUGUCA G UGGCGCUG	1307	CAGCGCCA GCCGAAAGGCGAGUCAAGGUCU UGACACAC	3220
1893	UGUCAGUG G CGCUGCCU	1308	AGGCAGCG GCCGAAAGGCGAGUCAAGGUCU CACUGACA	3221
1917	CUGCGCCA G CAGCAUGA	1309	UCAUGCUG GCCGAAAGGCGAGUCAAGGUCU UGGCGCAG	3222

Table 21

1920	CGCCAGCA G CAUGAUGA	1310	UCAUCAUG GCCGAAAGGCGAGUCAAGGUCU	UGCUGGCG	3223
1956	CUGCUGAA G UGAGGAGG	1311	CCUCCUCA GCCGAAAGGCGAGUCAAGGUCU	UUCAGCAG	3224
1964	GUGAGGAG G CCCAUGGG	1312	CCCAUGGG GCCGAAAGGCGAGUCAAGGUCU	CUCCUCAC	3225
1972	GCCCAUGG G CAGAAGAU	1313	AUCUUCUG GCCGAAAGGCGAGUCAAGGUCU	CCAUGGGC	3226
2006	ACACCUCC G UGUUUCAC	1314	GUGAACCA GCCGAAAGGCGAGUCAAGGUCU	GGAGGUGU	3227
2009	CCUCCGUG G UUCACUUU	1315	AAAGUGAA GCCGAAAGGCGAGUCAAGGUCU	CACGGAGG	3228
2019	UCACUUUG G UCACAAGU	1316	ACUUGUGA GCCGAAAGGCGAGUCAAGGUCU	CAAAGUGA	3229
2026	GGUCACAA G UAGGAGAC	1317	GUCUCCUA GCCGAAAGGCGAGUCAAGGUCU	UUGUGACC	3230
2042	CACAGAUG G CACCUGUG	1318	CACAGGUG GCCGAAAGGCGAGUCAAGGUCU	CAUCUGUG	3231
2051	CACCUGUG G CCAGAGCA	1319	UGCUCUGG GCCGAAAGGCGAGUCAAGGUCU	CACAGGUG	3232
2057	UGGCCAGA G CACCUCAG	1320	CUGAGGUG GCCGAAAGGCGAGUCAAGGUCU	UCUGGCCA	3233
2114	AGGAAAAG G CUGGCAAG	1321	CUUGCCAG GCCGAAAGGCGAGUCAAGGUCU	CUUUUCCU	3234
2118	AAAGGCUG G CAAGGUGG	1322	CCACCUUG GCCGAAAGGCGAGUCAAGGUCU	CAGCCUUU	3235
2123	CUGGCAAG G UGGGUUCC	1323	GGAACCCA GCCGAAAGGCGAGUCAAGGUCU	CUUGCCAG	3236
2127	CAAGGUGG G UUCCAGGG	1324	CCCUGGAA GCCGAAAGGCGAGUCAAGGUCU	CCACCUUG	3237
2172	AGAAAGAA G CACUCUGC	1325	GCAGAGUG GCCGAAAGGCGAGUCAAGGUCU	UUCUUUCU	3238
2183	CUCUGCUG G CGGGAUA	1326	UAUUCCCG GCCGAAAGGCGAGUCAAGGUCU	CAGCAGAG	3239
2198	UACUCUUG G UCACCUCA	1327	UGAGGUGA GCCGAAAGGCGAGUCAAGGUCU	CAAGAGUA	3240
2214	AAAUUUA G UCGGAAA	1328	UUUCCCGA GCCGAAAGGCGAGUCAAGGUCU	UUAAAUUU	3241
2243	AAACUUA G CCCUGAAC	1329	GUUCAGGG GCCGAAAGGCGAGUCAAGGUCU	UGAAGUUU	3242
2288	AACCCAAA G UAUUCUUC	1330	GAAGAAUA GCCGAAAGGCGAGUCAAGGUCU	UUUGGGUU	3243
2305	UUUUCUUA G UUUCAGAA	1331	UUCUGAAA GCCGAAAGGCGAGUCAAGGUCU	UAAGAAAA	3244
2314	UUUCAGAA G UACUGGCA	1332	UGCCAGUA GCCGAAAGGCGAGUCAAGGUCU	UUCUGAAA	3245
2320	AAGUACUG G CAUCACAC	1333	GUGUGAUG GCCGAAAGGCGAGUCAAGGUCU	CAGUACUU	3246
2333	ACACGCAG G UUACCUUG	1334	CAAGGUAA GCCGAAAGGCGAGUCAAGGUCU	CUGCGUGU	3247
2342	UUACCUUG G CGUGUGUC	1335	GACACACG GCCGAAAGGCGAGUCAAGGUCU	CAAGGUAA	3248
2344	ACCUUGGC G UGUGUCCC	1336	GGGACACA GCCGAAAGGCGAGUCAAGGUCU	GCCAAGGU	3249
2357	UCCCUGUG G UACCCUGG	1337	CCAGGGUA GCCGAAAGGCGAGUCAAGGUCU	CACAGGGA	3250
2365	GUACCCUG G CAGAGAAG	1338	CUUCUCUG GCCGAAAGGCGAGUCAAGGUCU	CAGGGUAC	3251
2381	GAGACCAA G CUUGUUUC	1339	GAAACAAG GCCGAAAGGCGAGUCAAGGUCU	UUGGUCUC	3252
2397	CCCUGCUG G CCAAAGUC	1340	GACUUUGG GCCGAAAGGCGAGUCAAGGUCU	CAGCAGGG	3253
2403	UGGCCAAA G UCAGUAGG	1341	CCUACUGA GCCGAAAGGCGAGUCAAGGUCU	UUUGGCCA	3254
2407	CAAAGUCA G UAGGAGAG	1342	CUCUCCUA GCCGAAAGGCGAGUCAAGGUCU	UGACUUUG	3255
2424	GAUGCACA G UUUGCUAU	1343	AUAGCAA GCCGAAAGGCGAGUCAAGGUCU	UGUGCAUC	3256
2463	AUAAACAA G CCUAACAU	1344	AUGUUAGG GCCGAAAGGCGAGUCAAGGUCU	UUGUUUAU	3257
2474	UAACAUTG G UGCAAAGA	1345	UCUUUGCA GCCGAAAGGCGAGUCAAGGUCU	CAAUGUUA	3258

Input Sequence = AF190725. Cut Site = G/.

Stem Length = 8. Core Sequence = GCcgaagGCGaGuCaaGGuCu

AF190725 (Homo sapiens beta-site APP cleaving enzyme (BACE) mRNA; 2526 bp)

Table 22

Table 22: Human BACE DNzyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
48	GCUGGAU A UGGUGGCC	3	GGCCACCA GGCTAGCTACAACGA AATCCAGC	3259
677	GCAGGGCU A CUACUGG	27	CCACGTAG GGCTAGCTACAACGA AGCCCTGC	3260
680	GGGCUACU A CGUGGAGA	28	TCTCCACG GGCTAGCTACAACGA AGTAGCCC	3261
733	UGGUGGAU A CAGGCAGC	31	GCTGCCTG GGCTAGCTACAACGA ATCCACCA	3262
788	GCAUCGCU A CUACCAGA	38	TCTGGTAG GGCTAGCTACAACGA AGCGATGC	3263
791	UCGCUACU A CCAGAGGC	39	GCCTCTGG GGCTAGCTACAACGA AGTAGCGA	3264
815	CAGCACAU A CCGGGACC	41	GGTCCCGG GGCTAGCTACAACGA ATGTGCTG	3265
839	GGGUGUGU A UGUGCCCU	43	AGGGCACA GGCTAGCTACAACGA ACACACCC	3266
848	UGUGCCCU A CACCCAGG	44	CCTGGGTG GGCTAGCTACAACGA AGGGCACA	3267
1004	GCUGGCCU A UGCUGAGA	58	TCTCAGCA GGCTAGCTACAACGA AGGCCAGC	3268
1171	UUGGAGGU A UCGACCAC	85	GTGGTGA GGCTAGCTACAACGA ACCTCAA	3269
1187	CUCGUGU A CACAGGCA	88	TGCTGTG GGCTAGCTACAACGA ACAGCGAG	3270
1205	UCUCUGGU A UACACCCA	91	TGGGTGTA GGCTAGCTACAACGA ACCAGAGA	3271
1207	UCUGGUAU A CACCCAUC	92	GATGGGTG GGCTAGCTACAACGA ATACCAGA	3272
1229	GGAGUGGU A UUAUGAGG	94	CCTCATAA GGCTAGCTACAACGA ACCACTCC	3273
1232	GUGGUAU A UGAGGUGA	96	TCACCTCA GGCTAGCTACAACGA AATACCAC	3274
1295	CAAGGAGU A CAACUAUG	101	CATAGTTG GGCTAGCTACAACGA ACTCCTTG	3275
1301	GUACAACU A UGACAAGA	102	TCTTGTC GGCTAGCTACAACGA AGTTGTAC	3276
1493	CUCACUCU A CCUAUUGG	130	CCATTAGG GGCTAGCTACAACGA AGAGTGAG	3277
1510	GUGAGGUU A CCAACCAG	133	CTGGTTGG GGCTAGCTACAACGA AACCTCAC	3278
1550	GCAGCAAU A CCUGCGGC	141	GCCGCAGG GGCTAGCTACAACGA ATTGCTGC	3279
1595	CGACUGUU A CAAGUUUG	144	CAAACCTG GGCTAGCTACAACGA AACAGTCG	3280
1633	GCACUGUU A UGGGAGCU	152	AGCTCCA GGCTAGCTACAACGA AACAGTGC	3281
1645	GAGCUGUU A UCAUGGAG	154	CTCCATGA GGCTAGCTACAACGA AACAGCTC	3282
1661	GGGCUUCU A CGUUGUCU	158	AGACAACG GGCTAGCTACAACGA AGAAGCCC	3283
1787	CUGUGGCU A CAACAUUC	176	GAATGTTG GGCTAGCTACAACGA AGCCACAG	3284
1832	CAUAGCCU A UGUCAUGG	182	CCATGACA GGCTAGCTACAACGA AGGCTATG	3285
2141	GGGACUGU A CCUGUAGG	212	CCTACAGG GGCTAGCTACAACGA ACAGTCCC	3286
2191	GCGGAAU A CUCUUGGU	215	ACCAAGAG GGCTAGCTACAACGA ATTCCCGC	3287
2290	CCCAAAGU A UUCUUCU	240	AAGAAGAA GGCTAGCTACAACGA ACTTTGGG	3288
2316	UCAGAAGU A CUGGCAUC	254	GATGCCAG GGCTAGCTACAACGA ACTTCTGA	3289
2336	CGCAGGUU A CCUUGGCG	257	CGCCAAGG GGCTAGCTACAACGA AACCTGCG	3290
2359	CCUGUGGU A CCCUGGCA	260	TGCCAGGG GGCTAGCTACAACGA ACCACAGG	3291
2431	AGUUUGCU A UUUGCUUU	269	AAAGCAA GGCTAGCTACAACGA AGCAAAC	3292
2455	GGGACUGU A UAAACAAG	275	CTTGTTTA GGCTAGCTACAACGA ACAGTCCC	3293
140	AAGCCGCC A CCGGCCCG	322	CGGGCCGG GGCTAGCTACAACGA GCGGCTT	3294
151	GGCCCGCC A UGCCCCCG	327	GCGGGGCA GGCTAGCTACAACGA GCGGGCC	3295
287	CGCUCUCC A CAGCCCGG	380	CCGGGCTG GGCTAGCTACAACGA GGAGAGCG	3296
368	GAGAAGCC A CCAGCACC	412	GGTGCTGG GGCTAGCTACAACGA GGCTTCTC	3297
374	CCACCAGC A CCACCCAG	415	CTGGGTGG GGCTAGCTACAACGA GCTGGTGG	3298
377	CCAGCACC A CCCAGACU	417	AGTCTGGG GGCTAGCTACAACGA GGTGCTGG	3299
451	CGGGGCC A CCAUGGCC	435	GGCCATGG GGCTAGCTACAACGA GGGCCCCG	3300
454	GGCCACC A UGGCCCAA	437	TTGGGCCA GGCTAGCTACAACGA GGTGGGCC	3301
512	GCCUGCCC A CGGCACCC	456	GGGTGCCG GGCTAGCTACAACGA GGGCAGGC	3302
517	CCCACGGC A CCCAGCAC	457	GTGCTGGG GGCTAGCTACAACGA GCCGTGGG	3303
524	CACCCAGC A CGGCAUCC	461	GGATGCCG GGCTAGCTACAACGA GCTGGGTG	3304
529	AGCACGGC A UCCGGCUG	462	CAGCCGGA GGCTAGCTACAACGA GCCGTGCT	3305

Table 22

721	CGCUCAAC A UCCUGGUG	508	CACCAGGA GGCTAGCTACAACGA GTTGAGCG	3306
770	UGCCCCC A CCCCUUCC	522	GGAAGGGG GGCTAGCTACAACGA GGGGGGCA	3307
782	CUUCCUGC A UCGCUACU	529	AGTAGCGA GGCTAGCTACAACGA GCAGGAAG	3308
811	UGUCCAGC A CAUACCGG	538	CCGGTATG GGCTAGCTACAACGA GCTGGACA	3309
813	UCCAGCAC A UACCGGGA	539	TCCCGGTA GGCTAGCTACAACGA GTGCTGGA	3310
850	UGCCCUAC A CCCAGGGC	547	GCCCTGGG GGCTAGCTACAACGA GTAGGGCA	3311
880	AGCUGGGC A CCGACCUG	553	CAGGTGGG GGCTAGCTACAACGA GCCCAGCT	3312
895	UGGUAAGC A UCCCCCAU	557	ATGGGGGA GGCTAGCTACAACGA GCTTACCA	3313
902	CAUCCCCC A UGGCCCCA	562	TGGGGCCA GGCTAGCTACAACGA GGGGGATG	3314
916	CCAACGUC A CUGUGCGU	567	ACGCACAG GGCTAGCTACAACGA GACGTTGG	3315
931	GUGCCAAC A UUGCUGCC	571	GGCAGCAA GGCTAGCTACAACGA GTTGGCAC	3316
940	UUGCUGCC A UCACUGAA	574	TTCAGTGA GGCTAGCTACAACGA GGCAGCAA	3317
943	CUGCCAUC A CUGAAUCA	575	TGATTGAG GGCTAGCTACAACGA GATGGCAG	3318
964	AGUUCUUC A UCAACGGC	580	GCCGTTGA GGCTAGCTACAACGA GAAGAACT	3319
988	GGGAAGGC A UCCUGGGG	586	CCCCAGGA GGCTAGCTACAACGA GCCTTCCC	3320
1070	GCAGACCC A CGUUCCCA	610	TGGGAACG GGCTAGCTACAACGA GGGTCTGC	3321
1156	GAGGGAGC A UGAUCAUU	638	AATGATCA GGCTAGCTACAACGA GCTCCCTC	3322
1162	GCAUGAUC A UUGGAGGU	639	ACCTCCAA GGCTAGCTACAACGA GATCATGC	3323
1178	UAUCGACC A CUCGUGU	641	ACAGCGAG GGCTAGCTACAACGA GGTCGATA	3324
1189	CGCUGUAC A CAGGCAGU	644	ACTGCCTG GGCTAGCTACAACGA GTACAGCG	3325
1209	UGGUUAUC A CCCAUCCG	649	CGGATGGG GGCTAGCTACAACGA GTATACCA	3326
1213	AUACACCC A UCCGGCGG	652	CCGCCGGA GGCTAGCTACAACGA GGGTGTAT	3327
1243	AGGUGAUC A UUGUGCGG	654	CCGCACAA GGCTAGCTACAACGA GATCACCT	3328
1312	ACAAGAGC A UUGUGGAC	663	GTCCACAA GGCTAGCTACAACGA GCTCTGT	3329
1327	ACAGUGGC A CCACCAAC	665	GTTGGTGG GGCTAGCTACAACGA GCCACTGT	3330
1330	GUGGCACC A CCAACCUU	667	AAGGTTGG GGCTAGCTACAACGA GGTGCCAC	3331
1378	UCAAUCC A UCAAGGCA	679	TGCCTTGA GGCTAGCTACAACGA GGATTGGA	3332
1396	CCUCCUCC A CGGAGAAG	687	CTTCTCCG GGCTAGCTACAACGA GGAGGAGG	3333
1456	AAGCAGGC A CCACCCCU	698	AGGGGTGG GGCTAGCTACAACGA GCCTGCTT	3334
1459	CAGGCACC A CCCCUUGG	700	CCAAGGGG GGCTAGCTACAACGA GGTGCCTG	3335
1471	CUUGGAAC A UUUUCCCA	705	TGGGAAAA GGCTAGCTACAACGA GTTCCAAG	3336
1483	UCCAGUC A UCUCACUC	709	GAGTGAGA GGCTAGCTACAACGA GACTGGGA	3337
1488	GUCAUCUC A CUCUACCU	711	AGGTAGAG GGCTAGCTACAACGA GAGATGAC	3338
1528	CCUUCGCG A UCACCAUC	723	GATGGTGA GGCTAGCTACAACGA GCGGAAGG	3339
1531	UCCGCAUC A CCAUCCUU	724	AAGGATGG GGCTAGCTACAACGA GATGCGGA	3340
1534	GCAUCACC A UCCUCCCG	726	CGGAAGGA GGCTAGCTACAACGA GGTGATGC	3341
1576	AUGUGGCC A CGUCCCAA	737	TTGGGACG GGCTAGCTACAACGA GGCCACAT	3342
1606	AGUUUGCC A UCUCACAG	744	CTGTGAGA GGCTAGCTACAACGA GGCAAAC	3343
1611	GCAUCUC A CAGUCAUC	746	GATGACTG GGCTAGCTACAACGA GAGATGGC	3344
1617	UCACAGUC A UCCACGGG	748	CCCGTGGA GGCTAGCTACAACGA GACTGTGA	3345
1621	AGUCAUCC A CGGGCACU	750	AGTGCCCG GGCTAGCTACAACGA GGATGACT	3346
1627	CCACGGGC A CUGUUAUG	751	CATAACAG GGCTAGCTACAACGA GCCCCTGG	3347
1648	CUGUUAUC A UGGAGGGC	754	GCCCTCCA GGCTAGCTACAACGA GATAACAG	3348
1715	CGCUUGCC A UGUGCACG	765	CGTGACAA GGCTAGCTACAACGA GGCAAGCG	3349
1721	CCAUGUGC A CGAUGAGU	766	ACTCATCG GGCTAGCTACAACGA GCACATGG	3350
1762	CUUUUGUC A CCUUGGAC	772	GTCCAAGG GGCTAGCTACAACGA GACAAAAG	3351
1771	CCUUGGAC A UGGAAGAC	775	GTCTTCCA GGCTAGCTACAACGA GTCCAAGG	3352
1792	GCUACAA A UUCCACAG	779	CTGTGGAA GGCTAGCTACAACGA GTTGTAGC	3353
1797	AACAUUCC A CAGACAGA	781	TCTGTCTG GGCTAGCTACAACGA GGAATGTT	3354
1819	CAACCCUC A UGACCAUA	788	TATGGTCA GGCTAGCTACAACGA GAGGGTTG	3355
1825	UCAUGACC A UAGCCUAU	790	ATAGGCTA GGCTAGCTACAACGA GGTGATGA	3356

Table 22

1837	CCU AUGUC A UGGCUGCC	793	GGCAGCCA GGCTAGCTACAACGA GACATAGG	3357
1846	UGGCUGCC A UCUGCGCC	796	GGCGCAGA GGCTAGCTACAACGA GGCAGCCA	3358
1861	CCCUCUUC A UGCUGCCA	802	TGGCAGCA GGCTAGCTACAACGA GAAGAGGG	3359
1869	AUGCUGCC A CUCUGCCU	805	AGGCAGAG GGCTAGCTACAACGA GGCAGCAT	3360
1879	UCUGCCUC A UGGUGUGU	810	ACACACCA GGCTAGCTACAACGA GAGGCAGA	3361
1922	CCAGCAGC A UGAUGACU	822	AGTCATCA GGCTAGCTACAACGA GCTGCTGG	3362
1942	CUGAUGAC A UCUCUCCUG	825	CAGGGAGA GGCTAGCTACAACGA GTCATCAG	3363
1968	GGAGGCC A UGGGCAGA	833	TCTGCCCA GGCTAGCTACAACGA GGGCCTCC	3364
1998	CCUGGACC A CACCUCCG	840	CGGAGGTG GGCTAGCTACAACGA GGTCCAGG	3365
2000	UGGACCAC A CCUCCGUG	841	CACGGAGG GGCTAGCTACAACGA GTGGTCCA	3366
2013	CGUGGUUC A CUUUGGUC	845	GACCAAAG GGCTAGCTACAACGA GAACCACG	3367
2022	CUUUGGUC A CAAGUAGG	847	CCTACTTG GGCTAGCTACAACGA GACCAAAG	3368
2035	UAGGAGAC A CAGAUGGC	849	GCCATCTG GGCTAGCTACAACGA GTCTCCTA	3369
2044	CAGAUGGC A CCUGUGGC	851	GCCACAGG GGCTAGCTACAACGA GCCATCTG	3370
2059	GCCAGAGC A CCUCAGGA	856	TCCTGAGG GGCTAGCTACAACGA GCTCTGGC	3371
2076	CCCUCUCC A CCCACCAA	866	TTGGTGGG GGCTAGCTACAACGA GGGGAGGG	3372
2080	CCCCACCC A CCAAUUGC	869	GCATTTGG GGCTAGCTACAACGA GGTGGGGG	3373
2174	AAAGAAGC A CUCUGCUG	885	CAGCAGAG GGCTAGCTACAACGA GCTTCTTT	3374
2201	UCUUGGUC A CCUCAAAU	891	ATTTGAGG GGCTAGCTACAACGA GACCAAGA	3375
2260	CUUUGUCC A CCAUUCU	906	AGGAATGG GGCTAGCTACAACGA GGACAAAG	3376
2263	UGUCCACC A UUCUUUA	908	TAAAGGAA GGCTAGCTACAACGA GGTGGACA	3377
2322	GUACUGGC A UCACACGC	922	GCGTGTGA GGCTAGCTACAACGA GCCAGTAC	3378
2325	CUGGCAUC A CACGCAGG	923	CCTGCGTG GGCTAGCTACAACGA GATGCCAG	3379
2327	GGCAUCAC A CGCAGGUU	924	AACCTGCG GGCTAGCTACAACGA GTGATGCC	3380
2421	GAGGAUGC A CAGUUGC	945	GCAAATCG GGCTAGCTACAACGA GCATCCTC	3381
2470	AGCCUAAC A UUGGUGCA	954	TGCACCAA GGCTAGCTACAACGA GTTAGGCT	3382
11	ACGCGUCC G CAGCCCGC	960	GCGGGCTG GGCTAGCTACAACGA GGACGCGT	3383
18	CGCAGCCC G CCGGGAG	961	CTCCCGGG GGCTAGCTACAACGA GGGCTGCG	3384
29	CGGGAGCU G CGAGCCGC	962	GCGGCTCG GGCTAGCTACAACGA AGCTCCCG	3385
36	UGCGAGCC G CGAGCUGG	964	CCAGCTCG GGCTAGCTACAACGA GGCTCGCA	3386
69	CAGCCAAC G CAGCCGCA	967	TGCGGCTG GGCTAGCTACAACGA GTTGCTG	3387
75	ACGCAGCC G CAGGAGCC	968	GGCTCCTG GGCTAGCTACAACGA GGCTGCGT	3388
94	GAGCCCUU G CCCUGCC	969	GGCAGGGG GGCTAGCTACAACGA AAGGGCTC	3389
100	UUGCCCUU G CCGCGGCC	970	GGCGCGGG GGCTAGCTACAACGA AGGGGCAA	3390
104	CCCUGCCC G CGCCGCCG	971	CGGCGGGG GGCTAGCTACAACGA GGGCAGGG	3391
106	CUGCCCGC G CCGCCGCC	972	GGCGGCGG GGCTAGCTACAACGA GCGGGCAG	3392
109	CCCGCGCC G CCGCCCGC	973	GCGGGCGG GGCTAGCTACAACGA GGCGCGGG	3393
112	GCGCCGCC G CCGCCCGG	974	CCGGCGGG GGCTAGCTACAACGA GGCGGCGC	3394
116	CGCCGCCG G CCGGGGGG	975	CCCCCGGG GGCTAGCTACAACGA GGGCGGGG	3395
137	GGGAAGCC G CCACCGGC	976	GCCGGTGG GGCTAGCTACAACGA GGCTTCCC	3396
148	ACCGGCCG G CCAUGCCC	977	GGGCATGG GGCTAGCTACAACGA GGGCCGGT	3397
153	CCCGCCAU G CCCGCCCC	978	GGGGCGGG GGCTAGCTACAACGA ATGGCGGG	3398
157	CCAUGCCC G CCCUCCC	979	GGGAGGGG GGCTAGCTACAACGA GGGCATGG	3399
172	CCAGCCCC G CCGGGAGC	980	GCTCCCGG GGCTAGCTACAACGA GGGGCTGG	3400
183	GGGAGCCC G CGCCCGCU	981	AGCGGGCG GGCTAGCTACAACGA GGGCTCCC	3401
185	GAGCCCGC G CCCGUGC	982	GCAGCGGG GGCTAGCTACAACGA GCGGGCTC	3402
189	CCGCGCCC G CUGCCAG	983	CTGGGCAG GGCTAGCTACAACGA GGGCGCGG	3403
192	CGCCCGCU G CCCAGGCU	984	AGCCTGGG GGCTAGCTACAACGA AGCGGGCG	3404
205	GGCUGGCC G CCGCCGUG	985	CACGGCGG GGCTAGCTACAACGA GGCCAGCC	3405
208	UGGCCGCC G CCGUGCCG	986	CGGCACGG GGCTAGCTACAACGA GGGGCCCA	3406
213	GCCGCCGU G CCGAUGUA	987	TACATCGG GGCTAGCTACAACGA ACGGCGGC	3407

Table 22

250	UCUCCCCU G CUCCCCGUG	989	CACGGGAG GGCTAGCTACAACGA AGGGGAGA	3408
258	GCUCCCCU G CUCUGCGG	990	CCGCAGAG GGCTAGCTACAACGA ACGGGAGC	3409
263	CGUGCUCU G CGGAUCUC	991	GAGATCCG GGCTAGCTACAACGA AGAGCACG	3410
280	CCCUGACC G CUCUCCAC	993	GTGGAGAG GGCTAGCTACAACGA GGTCAGGG	3411
320	AGGGCCCU G CAGGCCCU	994	AGGGCCTG GGCTAGCTACAACGA AGGGCCCT	3412
340	GUCCUGAU G CCCCCAAG	996	CTTGGGGG GGCTAGCTACAACGA ATCAGGAC	3413
397	GGGCAGGC G CCAGGGAC	998	GTCCCTGG GGCTAGCTACAACGA GCCTGCCC	3414
420	GGGCCAGU G CGAGCCCA	999	TGGGCTCG GGCTAGCTACAACGA ACTGGCCC	3415
468	CAAGCCCU G CCCUGGCU	1002	AGCCAGGG GGCTAGCTACAACGA AGGGCTTG	3416
480	UGGCUCCU G CUGUGGAU	1003	ATCCACAG GGCTAGCTACAACGA AGGAGCCA	3417
493	GGAUGGGC G CGGGAGUG	1004	CACTCCCG GGCTAGCTACAACGA GCCCATCC	3418
501	GCGGGAGU G CUGCCUGC	1005	GCAGGCAG GGCTAGCTACAACGA ACTCCCGC	3419
504	GGAGUGCU G CCUGCCCA	1006	TGGGCAGG GGCTAGCTACAACGA AGCACTCC	3420
508	UGCUGCCU G CCCACGGC	1007	GCCGTGGG GGCTAGCTACAACGA AGGCAGCA	3421
537	AUCCGGCU G CCCUGCG	1008	CGCAGGGG GGCTAGCTACAACGA AGCCGGAT	3422
543	CUGCCCCU G CGCAGCGG	1009	CCGCTGCG GGCTAGCTACAACGA AGGGGCAG	3423
545	GCCCCUGC G CAGCGGCC	1010	GGCGCTG GGCTAGCTACAACGA GCAGGGGC	3424
562	UGGGGGGC G CCCCCUG	1011	CAGGGGGG GGCTAGCTACAACGA GCCCCCA	3425
576	CUGGGGCU G CGGUGCC	1012	GGCAGCCG GGCTAGCTACAACGA AGCCCCAG	3426
582	CUGCGGCU G CCCCGGA	1013	TCCCGGGG GGCTAGCTACAACGA AGCCGAG	3427
708	AGCCCCC G CAGACGU	1019	AGCGTCTG GGCTAGCTACAACGA GGGGGCT	3428
714	CCGCAGAC G CUCAACAU	1020	ATGTTGAG GGCTAGCTACAACGA GTCTGCGG	3429
751	GUAACUUU G CAGUGGGU	1021	ACCCACTG GGCTAGCTACAACGA AAAGTTAC	3430
760	CAGUGGGU G CUGCCCCC	1022	GGGGGCAG GGCTAGCTACAACGA ACCCACTG	3431
763	UGGGUGCU G CCCCCAC	1023	GTGGGGGG GGCTAGCTACAACGA AGCACCCA	3432
780	CCCUUCCU G CAUCGCUA	1024	TAGCGATG GGCTAGCTACAACGA AGGAAGGG	3433
785	CCUGCAUC G CUACUACC	1025	GGTAGTAG GGCTAGCTACAACGA GATGCAGG	3434
843	GUGUAUGU G CCCUACAC	1026	GTGTAGGG GGCTAGCTACAACGA ACATACAC	3435
921	GUCACUGU G CGUGCCAA	1028	TTGCGACG GGCTAGCTACAACGA ACAGTGAC	3436
925	CUGUGCGU G CCAACAUU	1029	AATGTTGG GGCTAGCTACAACGA ACGCACAG	3437
934	CCAACAUU G CUGCCAUC	1030	GATGGCAG GGCTAGCTACAACGA AATGTTGG	3438
937	ACAUUGCU G CCAUCACU	1031	AGTGATGG GGCTAGCTACAACGA AGCAATGT	3439
1006	UGGCCUUAU G CUGAGAUU	1033	AATCTCAG GGCTAGCTACAACGA ATAGGCCA	3440
1015	CUGAGAUU G CCAGGCCU	1035	AGGCCTGG GGCTAGCTACAACGA AATCTCAG	3441
1092	UUCUCCCU G CAGCUUUG	1039	CAAAGCTG GGCTAGCTACAACGA AGGGAGAA	3442
1105	UUUGUGGU G CUGGCUUC	1040	GAAGCCAG GGCTAGCTACAACGA ACCACAAA	3443
1134	UCUGAAGU G CUGGCCUC	1042	GAGGCCAG GGCTAGCTACAACGA ACTTCAGA	3444
1182	GACCACUC G CUGUACAC	1045	GTGTACAG GGCTAGCTACAACGA GAGTGGTC	3445
1248	AUCAUUGU G CGGUGGGA	1048	TCCACCCG GGCTAGCTACAACGA ACAATGAT	3446
1286	AAUGGACU G CAAGGAGU	1050	ACTCCTTG GGCTAGCTACAACGA AGTCCATT	3447
1344	CUUCGUUU G CCCAAGAA	1052	TTCTTGGG GGCTAGCTACAACGA AAACGAAG	3448
1366	UUGAAGCU G CAGUCAAA	1054	TTTGACTG GGCTAGCTACAACGA AGCTTCAA	3449
1442	GCUGGUGU G CUGGCAAG	1056	CTTGCCAG GGCTAGCTACAACGA ACACCAGC	3450
1526	GUCCUUCG G CAUCACCA	1058	TGGTGATG GGCTAGCTACAACGA GGAAGGAC	3451
1542	AUCCUUCG G CAGCAUA	1059	TATTGCTG GGCTAGCTACAACGA GGAAGGAT	3452
1554	CAAUACCU G CGGCCAGU	1060	ACTGGCCG GGCTAGCTACAACGA AGGTATTG	3453
1603	ACAAGUUU G CCAUCUCA	1062	TGAGATGG GGCTAGCTACAACGA AAATCTGT	3454
1699	UUGGCUUU G CUGUCAGC	1066	GCTGACAG GGCTAGCTACAACGA AAAGCCAA	3455
1708	CUGUCAGC G CUUGCCAU	1067	ATGCAAG GGCTAGCTACAACGA GCTGACAG	3456
1712	CAGCGCUU G CCAUGUGC	1068	GCACATGG GGCTAGCTACAACGA AAGCGCTG	3457
1719	UGCCAUGU G CACGAUGA	1069	TCATCGTG GGCTAGCTACAACGA ACATGGCA	3458

Table 22

1843	UCAUGGCU G CCAUCUGC	1074	GCAGATGG GGCTAGCTACAACGA AGCCATGA	3459
1850	UGCCAUCU G CGCCUCUC	1075	AGAGGGCG GGCTAGCTACAACGA AGATGGCA	3460
1852	CCAUCUGC G CCCUCUUC	1076	GAAGAGGG GGCTAGCTACAACGA GCAGATGG	3461
1863	CUCUUAU G CUGCCACU	1077	AGTGGCAG GGCTAGCTACAACGA ATGAAGAG	3462
1866	UUCAUGCU G CCACUCUG	1078	CAGAGTGG GGCTAGCTACAACGA AGCATGAA	3463
1874	GCCACUCU G CCUCAUGG	1079	CCATGAGG GGCTAGCTACAACGA AGAGTGGC	3464
1895	UCAGUGGC G CUGCCUCC	1080	GGAGGCAG GGCTAGCTACAACGA GCCACTGA	3465
1898	GUGGCGCU G CCUCCGCU	1081	AGCGGAGG GGCTAGCTACAACGA AGCGCCAC	3466
1904	CUGCCUCC G CUGCCUGC	1082	GCAGGCAG GGCTAGCTACAACGA GGAGGCAG	3467
1907	CCUCCGCU G CCUGCGCC	1083	GGCGCAGG GGCTAGCTACAACGA AGCGGAGG	3468
1911	CGCUGCCU G CGCCAGCA	1084	TGCTGGCG GGCTAGCTACAACGA AGGCAGCG	3469
1913	CUGCCUGC G CCAGCAGC	1085	GCTGTGGG GGCTAGCTACAACGA GCAGGCAG	3470
1933	AUGACUUU G CUGAUGAC	1088	GTCATCAG GGCTAGCTACAACGA AAAGTCAT	3471
1950	AUCUCCCU G CUGAAGUG	1091	CACTTCAG GGCTAGCTACAACGA AGGGAGAT	3472
2087	CACCAAU G CCUCUGCC	1094	GGCAGAGG GGCTAGCTACAACGA ATTTGGTG	3473
2093	AUGCCUCU G CCUUGAUG	1095	CATCAAGG GGCTAGCTACAACGA AGAGGCAT	3474
2179	AGCACUCU G CUGGCGGG	1097	CCCGCCAG GGCTAGCTACAACGA AGAGTGCT	3475
2227	GAAAUUCU G CUGCUUGA	1098	TCAAGCAG GGCTAGCTACAACGA AGAATTTT	3476
2230	AUUCUGCU G CUUGAAAC	1099	GTTTCAAG GGCTAGCTACAACGA AGCAGAAT	3477
2329	CAUCACAC G CAGGUUAC	1102	GTAACCTG GGCTAGCTACAACGA GTGTGATG	3478
2393	GUUUCUUU G CUGGCCAA	1103	TTGGCCAG GGCTAGCTACAACGA AGGGAAAC	3479
2419	GAGAGGAU G CACAGUUU	1104	AAACTGTG GGCTAGCTACAACGA ATCCTCTC	3480
2428	CACAGUUU G CUUUUUGC	1105	GCAAATAG GGCTAGCTACAACGA AAATGTG	3481
2435	UGCUAUUU G CUUUAGAG	1106	CTCTAAAG GGCTAGCTACAACGA AAATGCA	3482
2476	ACAUUGGU G CAAAGAUU	1107	AATCTTTG GGCTAGCTACAACGA ACCAATGT	3483
2485	CAAAGAUU G CCUCUUGA	1108	TCAAGAGG GGCTAGCTACAACGA AATCTTTG	3484
219	GUGCCGAU G UAGCGGGC	1110	GCCCCTA GGCTAGCTACAACGA ATCGGCAC	3485
483	GUCUGCU G UGGAUGGG	1111	CCCATCCA GGCTAGCTACAACGA AGCAGGAG	3486
634	GCAGCUUU G UGGAGAUG	1112	CATCTCCA GGCTAGCTACAACGA AAAGCTGC	3487
804	AGGCAGCU G UCCAGCAC	1113	GTGCTGGA GGCTAGCTACAACGA AGCTGCCT	3488
835	GGAAGGGU G UGUUUGUG	1114	CACATACA GGCTAGCTACAACGA ACCCTTCC	3489
837	AAGGGUGU G UAUGUGCC	1115	GGCACATA GGCTAGCTACAACGA ACACCCTT	3490
841	GUGUGUAU G UGCCUAC	1116	GTAGGGCA GGCTAGCTACAACGA ATACACAC	3491
919	ACGUCACU G UGCGUGCC	1117	GGCACGCA GGCTAGCTACAACGA AGTGACGT	3492
1100	GCAGCUUU G UGGUGCUG	1118	CAGCACCA GGCTAGCTACAACGA AAAGCTGC	3493
1144	UGGCCUCU G UCGGAGGG	1119	CCCTCCGA GGCTAGCTACAACGA AGAGGCCA	3494
1185	CACUCGCU G UACACAGG	1120	CCTGTGTA GGCTAGCTACAACGA AGCGAGTG	3495
1246	UGAUCAUU G UGCGGGUG	1121	CACCGCA GGCTAGCTACAACGA AATGATCA	3496
1315	AGAGCAUU G UGGACAGU	1122	ACTGTCCA GGCTAGCTACAACGA AATGCTCT	3497
1356	AAGAAAGU G UUUGAAGC	1123	GCTTCAAA GGCTAGCTACAACGA ACTTTCTT	3498
1440	CAGCUGGU G UGUGGCA	1124	TGCCAGCA GGCTAGCTACAACGA ACCAGCTG	3499
1570	UGGAAGAU G UGGCCACG	1125	CGTGGCCA GGCTAGCTACAACGA ATCTTCCA	3500
1592	AGACGACU G UUAACAAGU	1126	ACTTGTA GGCTAGCTACAACGA AGTCGTCT	3501
1630	CGGGCACU G UUAUGGGA	1127	TCCCATAA GGCTAGCTACAACGA AGTGCCCG	3502
1642	UGGGAGCU G UUAUCAUG	1128	CATGATA GGCTAGCTACAACGA AGTCCCA	3503
1666	UCUACGUU G UCUUUGAU	1129	ATCAAAGA GGCTAGCTACAACGA AACGTAGA	3504
1702	GUUUUGCU G UCAGCGCU	1130	AGCGTGA GGCTAGCTACAACGA AGCAAAGC	3505
1717	CUUGCCAU G UGCACGAU	1131	ATCGTGCA GGCTAGCTACAACGA ATGGCAAG	3506
1759	GCCCUUUU G UCACCUUG	1132	CAAGGTGA GGCTAGCTACAACGA AAAAGGGC	3507
1781	GGAAGACU G UGGCUACA	1133	TGTAGCCA GGCTAGCTACAACGA AGTCTTCC	3508
1834	UAGCCUAU G UCAUGGCU	1134	AGCCATGA GGCTAGCTACAACGA ATAGGCTA	3509

Table 22

1884	CUCAUGGU G UGUCAGUG	1135	CACTGACA GGCTAGCTACAACGA ACCATGAG	3510
1886	CAUGGUGU G UCAGUGGC	1136	GCCACTGA GGCTAGCTACAACGA ACACCATG	3511
2048	UGGCACCU G UGGCCAGA	1137	TCTGGCCA GGCTAGCTACAACGA AGGTGCCA	3512
2139	CAGGGACU G UACCUGUA	1138	TACAGGTA GGCTAGCTACAACGA AGTCCTTG	3513
2145	CUGUACCU G UAGGAAAC	1139	GTTTCCTA GGCTAGCTACAACGA AGGTACAG	3514
2256	GAACCUUU G UCCACCAU	1140	ATGGTGGA GGCTAGCTACAACGA AAAGGTTC	3515
2346	CUUGGCGU G UGUCCUG	1141	CAGGGACA GGCTAGCTACAACGA ACGCCAAG	3516
2348	UGGCGUGU G UCCUGUG	1142	CACAGGGA GGCTAGCTACAACGA ACACGCCA	3517
2354	GUGUCCCU G UGUACCC	1143	GGGTACCA GGCTAGCTACAACGA AGGGACAC	3518
2385	CCAAGCUU G UUUCCUG	1144	CAGGGAAA GGCTAGCTACAACGA AAGCTTGG	3519
2453	CAGGGACU G UAUAAACA	1145	TGTTTATA GGCTAGCTACAACGA AGTCCCTG	3520
14	CGUCCGCA G CCCGCCG	1146	CGGGCGGG GGCTAGCTACAACGA TGCGGACG	3521
26	GCCCGGGA G CUGCGAGC	1147	GCTCGCAG GGCTAGCTACAACGA TCCCGGGC	3522
33	AGCUGCGA G CCGCGAGC	1148	GCTCGCGG GGCTAGCTACAACGA TCGCAGCT	3523
40	AGCCGCGA G CUGGAUUA	1149	TAATCCAG GGCTAGCTACAACGA TCGCGGCT	3524
51	GGAUUAUG G UGGCCUGA	1150	TCAGGCCA GGCTAGCTACAACGA CATAATCC	3525
54	UUAUGGUG G CCUGAGCA	1151	TGCTCAGG GGCTAGCTACAACGA CACCATAA	3526
60	UGGCCUGA G CAGCCAAC	1152	GTTGGCTG GGCTAGCTACAACGA TCAGGCCA	3527
63	CCUGAGCA G CCAACGCA	1153	TGCGTTGG GGCTAGCTACAACGA TGCTCAGG	3528
72	CCAACGCA G CCGCAGGA	1154	TCCTGCGG GGCTAGCTACAACGA TCGGTTGG	3529
81	CCGCAGGA G CCCGGAGC	1155	GCTCCGGG GGCTAGCTACAACGA TCCTGCGG	3530
88	AGCCCGGA G CCCUUGCC	1156	GGCAAGGG GGCTAGCTACAACGA TCCGGGCT	3531
134	CCAGGGAA G CCGCCACC	1157	GGTGCGGG GGCTAGCTACAACGA TTCCCTGG	3532
144	CGCCACCG G CCCGCCAU	1158	ATGGCGGG GGCTAGCTACAACGA CGGTGGCG	3533
167	CCUCCCA G CCCGCGG	1159	CGGCGGGG GGCTAGCTACAACGA TGGGAGGG	3534
179	CGCCGGGA G CCCGCGC	1160	GGCGCGGG GGCTAGCTACAACGA TCCCGGCG	3535
198	CGCCCCAG G CUGGCCGC	1161	GCGGCCAG GGCTAGCTACAACGA CTGGGCAG	3536
202	CCAGGCUG G CCGCCGCC	1162	GGCGGCGG GGCTAGCTACAACGA CAGCCTGG	3537
211	CCGCCGCC G UGCCGAUG	1163	CATCGGCA GGCTAGCTACAACGA GCGGCGGG	3538
222	CCGAUGUA G CGGGCUCC	1164	GGAGCCCG GGCTAGCTACAACGA TACATCGG	3539
226	UGUAGCGG G CUCCGAU	1165	ATCCGGAG GGCTAGCTACAACGA CCGCTACA	3540
239	GAUCCCA G CCUCUCCC	1166	GGGAGAGG GGCTAGCTACAACGA TGGGATCC	3541
256	CUGCUCCC G UGCUUGC	1167	GCAGAGCA GGCTAGCTACAACGA GGGAGCAG	3542
290	UCUCCACA G CCCGGACC	1168	GGTCCGGG GGCTAGCTACAACGA TGTGGAGA	3543
304	ACCCGGGG G CUGGCCCA	1169	TGGGCCAG GGCTAGCTACAACGA CCCCGGGT	3544
308	GGGGGCGU G CCCAGGGC	1170	GCCCTGGG GGCTAGCTACAACGA CAGCCCCC	3545
315	GGCCCAGG G CCCUGCAG	1171	CTGCAGGG GGCTAGCTACAACGA CCTGGGCC	3546
324	CCCUGCAG G CCCUGGCG	1172	CGCCAGGG GGCTAGCTACAACGA CTGCAGGG	3547
330	AGGCCUG G CGUCCUGA	1173	TCAGGACG GGCTAGCTACAACGA CAGGGCCT	3548
332	GCCCUGGC G UCCUGAUG	1174	CATCAGGA GGCTAGCTACAACGA GCCAGGGC	3549
348	GCCCCCAA G CUCCUCU	1175	AGAGGGAG GGCTAGCTACAACGA TTGGGGGC	3550
365	CCUGAGAA G CCACCAGC	1176	GCTGGTGG GGCTAGCTACAACGA TTCTCAGG	3551
372	AGCCACCA G CACCACCC	1177	GGGTGGTG GGCTAGCTACAACGA TGGTGGCT	3552
391	ACUUGGGG G CAGGCGCC	1178	GGCGCCTG GGCTAGCTACAACGA CCCCAAGT	3553
395	GGGGGCGA G CGCCAGGG	1179	CCCTGGCG GGCTAGCTACAACGA CTGCCCCC	3554
410	GGACGGAC G UGGGCCAG	1180	CTGGCCCA GGCTAGCTACAACGA GTCCGTCC	3555
414	GGACGUGG G CCAGUGCG	1181	CGCACTGG GGCTAGCTACAACGA CCACGTCC	3556
418	GUGGGCCA G UGCGAGCC	1182	GGCTCGCA GGCTAGCTACAACGA TGGCCAC	3557
424	CAGUGCGA G CCCAGAGG	1183	CCTCTGGG GGCTAGCTACAACGA TCGACTG	3558
433	CCCAGAGG G CCCGAAGG	1184	CCTTCGGG GGCTAGCTACAACGA CCTCTGGG	3559
441	GCCCAAG G CCGGGGCC	1185	GGCCCCGG GGCTAGCTACAACGA CTTGGGGC	3560

Table 22

447	AGGCCGGG G CCCACCAU	1186	ATGGTGGG GGCTAGCTACAACGA CCCGGCCT	3561
457	CCACCAUG G CCCAAGCC	1187	GGCTTGGG GGCTAGCTACAACGA CATGGTGG	3562
463	UGGCCCAA G CCCUGCCC	1188	GGGCAGGG GGCTAGCTACAACGA TTGGGCCA	3563
474	CUGCCUG G CUCCUGCU	1189	AGCAGGAG GGCTAGCTACAACGA CAGGGCAG	3564
491	GUGGAUGG G CGCGGGAG	1190	CTCCCGCG GGCTAGCTACAACGA CCATCCAC	3565
499	GCGCGGGA G UGUGCCU	1191	AGGCAGCA GGCTAGCTACAACGA TCCCGCGC	3566
515	UGCCACG G CACCCAGC	1192	GCTGGGTG GGCTAGCTACAACGA CGTGGGCA	3567
522	GGCACCA G CACGGCAU	1193	ATGCCGTG GGCTAGCTACAACGA TGGGTGCC	3568
527	CCAGCAG G CAUCCGGC	1194	GCCGGATG GGCTAGCTACAACGA CGTGCTGG	3569
534	GGCAUCCG G CUGCCCCU	1195	AGGGGCAG GGCTAGCTACAACGA CGGATGCC	3570
548	CCUGCGCA G CGGCCUGG	1196	CCAGGCCG GGCTAGCTACAACGA TGCGCAGG	3571
551	GCGCAGCG G CUUGGGGG	1197	CCCCCAGG GGCTAGCTACAACGA CGTGCGCG	3572
560	CCUGGGGG G CGCCCCC	1198	GGGGGGCG GGCTAGCTACAACGA CCCCCAGG	3573
573	CCCCUGGG G CUGCGGCU	1199	AGCCGCAG GGCTAGCTACAACGA CCCAGGGG	3574
579	GGGUGCG G CUGCCCCG	1200	CGGGGCAG GGCTAGCTACAACGA CGCAGCCC	3575
603	GACGAAGA G CCCGAGGA	1201	TCCTCGGG GGCTAGCTACAACGA TCTTCGTC	3576
612	CCCGAGGA G CCCGGCCG	1202	CGGCCGGG GGCTAGCTACAACGA TCCTCGGG	3577
617	GGAGCCCG G CCGGAGGG	1203	CCCTCCGG GGCTAGCTACAACGA CGGGCTCC	3578
626	CCGGAGGG G CAGCUUUG	1204	CAAAGCTG GGCTAGCTACAACGA CCCTCCGG	3579
629	GAGGGGCA G CUUUGUGG	1205	CCACAAAG GGCTAGCTACAACGA TGCCCTCT	3580
643	UGGAGAUG G UGGACAAC	1206	GTTGTCCA GGCTAGCTACAACGA CATCTCCA	3581
659	CCUGAGGG G CAAGUCGG	1207	CCGACTTG GGCTAGCTACAACGA CCCTCAGG	3582
663	AGGGGCAA G UCGGGGCA	1208	TGCCCCGA GGCTAGCTACAACGA TTGCCCTT	3583
669	AAGUCGGG G CAGGGCUA	1209	TAGCCCTG GGCTAGCTACAACGA CCCGACTT	3584
674	GGGGCAGG G CUACUACG	1210	CGTAGTAG GGCTAGCTACAACGA CCTGCCCC	3585
682	GUACUAC G UGGAGAUG	1211	CATCTCCA GGCTAGCTACAACGA GTAGTAGC	3586
694	AGAUGACC G UGGGCAGC	1212	GCTGCCCA GGCTAGCTACAACGA GGTCATCT	3587
698	GACCGUGG G CAGCCCCC	1213	GGGGGCTG GGCTAGCTACAACGA CCACGGTC	3588
701	CGUGGGCA G CCCCCCGC	1214	GCGGGGGG GGCTAGCTACAACGA TGCCCACG	3589
727	ACAUCCUG G UGGAUACA	1215	TGTATCCA GGCTAGCTACAACGA CAGGATGT	3590
737	GGAUACAG G CAGCAGUA	1216	TACTGCTG GGCTAGCTACAACGA CTGTATCC	3591
740	UACAGGCA G CAGUAACU	1217	AGTTACTG GGCTAGCTACAACGA TGCCTGTA	3592
743	AGGCAGCA G UAACUUG	1218	CAAAGTTA GGCTAGCTACAACGA TGCTGCCT	3593
754	ACUUGCA G UGGGUGCU	1219	AGCACCCA GGCTAGCTACAACGA TGCAAAGT	3594
758	UGCAGUGG G UGUGCCC	1220	GGGCAGCA GGCTAGCTACAACGA CCACTGCA	3595
798	UACCAGAG G CAGCUGUC	1221	GACAGCTG GGCTAGCTACAACGA CTCTGGTA	3596
801	CAGAGGCA G CUGUCCAG	1222	CTGGACAG GGCTAGCTACAACGA TGCCTCTG	3597
809	GCUGUCCA G CACAUACC	1223	GGTATGTG GGCTAGCTACAACGA TGGACAGC	3598
833	CCGGAAGG G UGUGUAUG	1224	CATACACA GGCTAGCTACAACGA CCTTCCGG	3599
857	CACCCAGG G CAAGUGGG	1225	CCCCTTGG GGCTAGCTACAACGA CCTGGGTG	3600
861	CAGGGCAA G UGGGAAGG	1226	CCTTCCCA GGCTAGCTACAACGA TTGCCCTG	3601
873	GAAGGGGA G CUGGGCAC	1227	GTGCCCAG GGCTAGCTACAACGA TCCCTTTC	3602
878	GGAGCUGG G CACCGACC	1228	GGTCGGTG GGCTAGCTACAACGA CCAGCTCC	3603
889	CCGACCUG G UAAGCAUC	1229	GATGCTTA GGCTAGCTACAACGA CAGGTCGG	3604
893	CCUGGUAA G CAUCCCCC	1230	GGGGGATG GGCTAGCTACAACGA TTACCAGG	3605
905	CCCCAUG G CCCCAACG	1231	CGTTGGGG GGCTAGCTACAACGA CATGGGGG	3606
913	GCCCCAAC G UCACUGUG	1232	CACAGTGA GGCTAGCTACAACGA GTTGGGGC	3607
923	CACUGUGC G UGCCAACA	1233	TGTTGGCA GGCTAGCTACAACGA GCACAGTG	3608
957	UCAGACAA G UUCUUCAU	1234	ATGAAGAA GGCTAGCTACAACGA TTGTCTGA	3609
971	CAUCAACG G CUCCAACU	1235	AGTTGGAG GGCTAGCTACAACGA CGTTGATG	3610
986	CUGGGAAG G CAUCCUGG	1236	CCAGGATG GGCTAGCTACAACGA CTTCCAG	3611

Table 22

996	AUCCUGGG G CUGGCCUA	1237	TAGGCCAG GGCTAGCTACAACGA CCCAGGAT	3612
1000	UGGGGUG G CCUAUGCU	1238	AGCATAGG GGCTAGCTACAACGA CAGCCCCA	3613
1020	AUUGCCAG G CCUGACGA	1239	TCGTCAGG GGCTAGCTACAACGA CTGGCAAT	3614
1038	UCCCUGGA G CCUUUCUU	1240	AAGAAAGG GGCTAGCTACAACGA TCCAGGGA	3615
1057	ACUCUCUG G UAAAGCAG	1241	CTGCTTTA GGCTAGCTACAACGA CAGAGAGT	3616
1062	CUGGUAAA G CAGACCCA	1242	TGGGTCTG GGCTAGCTACAACGA TTTACCAG	3617
1072	AGACCCAC G UUCCCAAC	1243	GTTGGGAA GGCTAGCTACAACGA GTGGGTCT	3618
1095	UCCCUGCA G CUUUGUGG	1244	CCACAAAG GGCTAGCTACAACGA TGCAGGGA	3619
1103	GCUUUGUG G UGUGGCU	1245	AGCCAGCA GGCTAGCTACAACGA CACAAAGC	3620
1109	UGGUGCUG G CUUCCCCC	1246	GGGGGAAG GGCTAGCTACAACGA CAGCACCA	3621
1125	CUCAACCA G UCUGAAGU	1247	ACTTCAGA GGCTAGCTACAACGA TGGTTGAG	3622
1132	AGUCUGAA G UGUGGCC	1248	GGCCAGCA GGCTAGCTACAACGA TTCAGACT	3623
1138	AAGUGCUG G CCUCUGUC	1249	GACAGAGG GGCTAGCTACAACGA CAGCACTT	3624
1154	CGGAGGGA G CAUGAUCA	1250	TGATCATG GGCTAGCTACAACGA TCCCTCCG	3625
1169	CAUUGGAG G UAUCGACC	1251	GGTCGATA GGCTAGCTACAACGA CTCCAATG	3626
1193	GUACACAG G CAGUCUCU	1252	AGAGACTG GGCTAGCTACAACGA CTGTGTAC	3627
1196	CACAGGCA G UCUCUGGU	1253	ACCAGAGA GGCTAGCTACAACGA TGCCTGTG	3628
1203	AGUCUCUG G UAUACACC	1254	GGTGATA GGCTAGCTACAACGA CAGAGACT	3629
1218	CCCAUCCG G CGGGAGUG	1255	CACTCCCG GGCTAGCTACAACGA CGGATGGG	3630
1224	CGGCGGGA G UGUUAUUA	1256	TAATACCA GGCTAGCTACAACGA TCCGCGCG	3631
1227	CGGGAGUG G UAUUAUGA	1257	TCATAATA GGCTAGCTACAACGA CACTCCCG	3632
1237	AUUAUGAG G UGAUCAUU	1258	AATGATCA GGCTAGCTACAACGA CTCATAAT	3633
1252	UUGUGCGG G UGGAGAUC	1259	GATCTCCA GGCTAGCTACAACGA CCGCACAA	3634
1293	UGCAAGGA G UACAACUA	1260	TAGTTGTA GGCTAGCTACAACGA TCCTTGCA	3635
1310	UGACAAGA G CAUUGUGG	1261	CCACAATG GGCTAGCTACAACGA TCTTGTC	3636
1322	UGUGGACA G UGGCACCA	1262	TGGTGCCA GGCTAGCTACAACGA TGTCCACA	3637
1325	GGACAGUG G CACCACCA	1263	TGGTGGTG GGCTAGCTACAACGA CACTGTCC	3638
1340	CAACCUUC G UUUGCCA	1264	TGGGCAA GGCTAGCTACAACGA GAAGGTTG	3639
1354	CCAAGAAA G UGUUUGAA	1265	TTCAAACA GGCTAGCTACAACGA TTTCTTGG	3640
1363	UGUUUGAA G CUGCAGUC	1266	GACTGCAG GGCTAGCTACAACGA TTCAAACA	3641
1369	AAGUGCA G UCAAAUCC	1267	GGATTGTA GGCTAGCTACAACGA TGCAGCTT	3642
1384	CCAUCAAG G CAGCCUCC	1268	GGAGGCTG GGCTAGCTACAACGA CTTGATGG	3643
1387	UCAAGGCA G CCUCCUCC	1269	GGAGGAGG GGCTAGCTACAACGA TGCCTTGA	3644
1404	ACGAGAA G UUCCUGA	1270	TCAGGGAA GGCTAGCTACAACGA TTCTCCGT	3645
1415	CCUGAUG G UUUCUGG	1271	GCCAGAAA GGCTAGCTACAACGA CATCAGGG	3646
1422	GGUUCUG G CUAGGAGA	1272	TCTCCTAG GGCTAGCTACAACGA CAGAAACC	3647
1431	CUAGGAGA G CAGCUGGU	1273	ACCAGCTG GGCTAGCTACAACGA TCTCCTAG	3648
1434	GGAGAGCA G CUGGUGUG	1274	CACACCAG GGCTAGCTACAACGA TGCTCTCC	3649
1438	AGCAGCUG G UGUGCUGG	1275	CCAGCACA GGCTAGCTACAACGA CAGCTGCT	3650
1446	GUGUGCUG G CAAGCAGG	1276	CCTGCTTG GGCTAGCTACAACGA CAGCACAC	3651
1450	GCUGGCAA G CAGGCACC	1277	GGTGCTTG GGCTAGCTACAACGA TTGCCAGC	3652
1454	GCAAGCAG G CACCACCC	1278	GGGTGGTG GGCTAGCTACAACGA CTGCTTGC	3653
1480	UUUUCCCA G UCAUCUCA	1279	TGAGATGA GGCTAGCTACAACGA TGGGAAAA	3654
1502	CCUAAUGG G UGAGGUUA	1280	TAACCTCA GGCTAGCTACAACGA CCATTAGG	3655
1507	UGGGUGAG G UUACCAAC	1281	GTTGGTAA GGCTAGCTACAACGA CTCACCCA	3656
1518	ACCAACCA G UCCUCCG	1282	CGGAAGGA GGCTAGCTACAACGA TGGTTGGT	3657
1545	CUUCCGCA G CAAUACCU	1283	AGGTATTG GGCTAGCTACAACGA TGCGGAAG	3658
1557	UACCGCG G CCAGUGGA	1284	TCCACTGG GGCTAGCTACAACGA CGCAGGTA	3659
1561	UGCGGCCA G UGGAAGAU	1285	ATCTTCCA GGCTAGCTACAACGA TGGCCGCA	3660
1573	AAGAUGUG G CCACGUCC	1286	GGACGTGG GGCTAGCTACAACGA CACATCTT	3661
1578	GUGGCCAC G UCCCAAGA	1287	TCTTGGGA GGCTAGCTACAACGA GTGGCCAC	3662

Table 22

1599	UGUJACAA G UUUGCCAU	1288	ATGGCAAA GGCTAGCTACAACGA TTGTAACA	3663
1614	AUCUCACA G UCAUCCAC	1289	GTGGATGA GGCTAGCTACAACGA TGTGAGAT	3664
1625	AUCCACGG G CACUGUUA	1290	TAACAGTG GGCTAGCTACAACGA CCGTGGAT	3665
1639	UUAUGGGA G CUGUUAUC	1291	GATAACAG GGCTAGCTACAACGA TCCCATAA	3666
1655	CAUGGAGG G CUUCUACG	1292	CGTAGAAG GGCTAGCTACAACGA CCTCCATG	3667
1663	GCUUCUAC G UUGUCUUU	1293	AAAGACAA GGCTAGCTACAACGA GTAGAAGC	3668
1678	UUGAUCGG G CCCGAAAA	1294	TTTTCGGG GGCTAGCTACAACGA CCGATCAA	3669
1694	ACGAUUG G CUUUGCUG	1295	CAGCAAAG GGCTAGCTACAACGA CAATTTCGT	3670
1706	UGCUGUCA G CGCUUGCC	1296	GGCAAGCG GGCTAGCTACAACGA TGACAGCA	3671
1728	CACGAUGA G UUCAGGAC	1297	GTCTGAA GGCTAGCTACAACGA TCATCGTG	3672
1738	UCAGGACG G CAGCGGUG	1298	CACCGCTG GGCTAGCTACAACGA CGTCCTGA	3673
1741	GGACGGCA G CGGUGGAA	1299	TTCCACCG GGCTAGCTACAACGA TGCCGTCC	3674
1744	CGGCAGCG G UGGAAGGC	1300	GCCTTCCA GGCTAGCTACAACGA CGCTGCCG	3675
1751	GGUGGAAG G CCCUUUUG	1301	CAAAAGGG GGCTAGCTACAACGA CTTCCACC	3676
1784	AGACUGUG G CUACAACA	1302	TGTTGTAG GGCTAGCTACAACGA CACAGTCT	3677
1809	ACAGAUGA G UCAACCCU	1303	AGGGTTGA GGCTAGCTACAACGA TCATCTGT	3678
1828	UGACCAUA G CCUAUGUC	1304	GACATAGG GGCTAGCTACAACGA TATGGTCA	3679
1840	AUGUCAUG G CUGCCAUC	1305	GATGGCAG GGCTAGCTACAACGA CATGACAT	3680
1882	GCCUCAUG G UGUGUCAG	1306	CTGACACA GGCTAGCTACAACGA CATGAGGC	3681
1890	GUGUGUCA G UGGCGCUG	1307	CAGCGCCA GGCTAGCTACAACGA TGACACAC	3682
1893	UGUCAGUG G CGCUGCCU	1308	AGGCAGCG GGCTAGCTACAACGA CACTGACA	3683
1917	CUGCGCCA G CAGCAUGA	1309	TCATGCTG GGCTAGCTACAACGA TGGCGCAG	3684
1920	CGCCAGCA G CAUGAUGA	1310	TCATCATG GGCTAGCTACAACGA TGCTGGCG	3685
1956	CUGCUGAA G UGAGGAGG	1311	CCTCTCA GGCTAGCTACAACGA TTCAGCAG	3686
1964	GUGAGGAG G CCCAUGGG	1312	CCCATGGG GGCTAGCTACAACGA CTCCTCAC	3687
1972	GCCCAUGG G CAGAAGAU	1313	ATCTTCTG GGCTAGCTACAACGA CCATGGGC	3688
2006	ACACCUCC G UGUUUCAC	1314	GTGAACCA GGCTAGCTACAACGA GGAGGTGT	3689
2009	CCUCCGUG G UUCACUUU	1315	AAAGTGAA GGCTAGCTACAACGA CACGGAGG	3690
2019	UCACUUUG G UCACAAGU	1316	ACTTGTGA GGCTAGCTACAACGA CAAAGTGA	3691
2026	GGUCACAA G UAGGAGAC	1317	GTCTCCTA GGCTAGCTACAACGA TTGTGACC	3692
2042	CACAGAUG G CACCUGUG	1318	CACAGGTG GGCTAGCTACAACGA CATCTGTG	3693
2051	CACCUGUG G CCAGAGCA	1319	TGCTCTGG GGCTAGCTACAACGA CACAGGTG	3694
2057	UGGCCAGA G CACCUCAG	1320	CTGAGGTG GGCTAGCTACAACGA TCTGGCCA	3695
2114	AGGAAAAG G CUGGCAAG	1321	CTTGCCAG GGCTAGCTACAACGA CTTTTCCT	3696
2118	AAAGGCUG G CAAGGUGG	1322	CCACCTTG GGCTAGCTACAACGA CAGCCTTT	3697
2123	CUGGCAAG G UGGGUUCC	1323	GGAACCCA GGCTAGCTACAACGA CTTGCCAG	3698
2127	CAAGGUGG G UUCAGGG	1324	CCCTGGAA GGCTAGCTACAACGA CCACCTTG	3699
2172	AGAAAGAA G CACUCUGC	1325	GCAGAGTG GGCTAGCTACAACGA TTCTTTCT	3700
2183	CUCUGCUG G CGGGAUA	1326	TATTCCTG GGCTAGCTACAACGA CAGCAGAG	3701
2198	UACUCUUG G UCACCUCA	1327	TGAGGTGA GGCTAGCTACAACGA CAAGAGTA	3702
2214	AAAUUUA G UCGGAAA	1328	TTTCCCGA GGCTAGCTACAACGA TTAAATTT	3703
2243	AAACUUA G CCCUGAAC	1329	GTTACGGG GGCTAGCTACAACGA TGAAGTTT	3704
2288	AACCCAAA G UAUUCUUC	1330	GAAGAATA GGCTAGCTACAACGA TTTGGGTT	3705
2305	UUUUCUUA G UUUCAGAA	1331	TTCTGAAA GGCTAGCTACAACGA TAAGAAAA	3706
2314	UUUCAGAA G UACUGGCA	1332	TGCCAGTA GGCTAGCTACAACGA TTCTGAAA	3707
2320	AAGUACUG G CAUCACAC	1333	GTGTGATG GGCTAGCTACAACGA CAGTACTT	3708
2333	ACACGCAG G UUACCUUG	1334	CAAGGTAA GGCTAGCTACAACGA CTGCGTGT	3709
2342	UUACCUUG G CGUGUGUC	1335	GACACACG GGCTAGCTACAACGA CAAGGTAA	3710
2344	ACCUUGGC G UGUGUCCC	1336	GGGACACA GGCTAGCTACAACGA GCCAAGGT	3711
2357	UCCCUUG G UACCCUGG	1337	CCAGGGTA GGCTAGCTACAACGA CACAGGGA	3712
2365	GUACCCUG G CAGAGAAG	1338	CTTCTCTG GGCTAGCTACAACGA CAGGGTAC	3713

Table 22

2381	GAGACCAA G CUUGUUUC	1339	GAAACAAG GGCTAGCTACAACGA TTGGTCTC	3714
2397	CCCUGCUG G CCAAAGUC	1340	GACTTTGG GGCTAGCTACAACGA CAGCAGGG	3715
2403	UGGCCAAA G UCAGUAGG	1341	CCTACTGA GGCTAGCTACAACGA TTTGGCCA	3716
2407	CAAAGUCA G UAGGAGAG	1342	CTCTCCTA GGCTAGCTACAACGA TGACTTTG	3717
2424	GAUGCACA G UUUGCUAU	1343	ATAGCAAA GGCTAGCTACAACGA TGTGCATC	3718
2463	AUAAACAA G CCUAACAU	1344	ATGTTAGG GGCTAGCTACAACGA TTGTTTAT	3719
2474	UAACAUUG G UGCAAGA	1345	TCTTTGCA GGCTAGCTACAACGA CAATGTTA	3720
45	CGAGCUGG A UUAUGGUG	1346	CACCATAA GGCTAGCTACAACGA CCAGCTCG	3721
67	AGCAGCCA A CGCAGCCG	1347	CGGCTGCG GGCTAGCTACAACGA TGGCTGCT	3722
125	CCGGGGGG A CCAGGGAA	1348	TTCCCTGG GGCTAGCTACAACGA CCCCCCGG	3723
217	CCGUGCCG A UGUAGCGG	1349	CCGCTACA GGCTAGCTACAACGA CGGCACGG	3724
233	GGCUCGG A UCCCAGCC	1350	GGCTGGGA GGCTAGCTACAACGA CCGGAGCC	3725
267	CUCUGCGG A UCUCCCCU	1351	AGGGGAGA GGCTAGCTACAACGA CCGCAGAG	3726
277	CUCCCCUG A CCGCUCUC	1352	GAGAGCGG GGCTAGCTACAACGA CAGGGGAG	3727
296	CAGCCCGG A CCCGGGGG	1353	CCCCCGGG GGCTAGCTACAACGA CCGGGCTG	3728
338	GCGUCCG A UGCCCCA	1354	TGGGGGCA GGCTAGCTACAACGA CAGGACGC	3729
383	CCACCCAG A CUUGGGGG	1355	CCCCAAG GGCTAGCTACAACGA CTGGGTGG	3730
404	CGCCAGGG A CGGACGUG	1356	CACGTCGG GGCTAGCTACAACGA CCTGGCG	3731
408	AGGGACGG A CGUGGGCC	1357	GGCCACG GGCTAGCTACAACGA CCGTCCCT	3732
487	UGCUGUGG A UGGGCGCG	1358	CGCGCCA GGCTAGCTACAACGA CCACAGCA	3733
592	CCCGGGAG A CCGACGAA	1359	TTCGTCGG GGCTAGCTACAACGA CTCCCGGG	3734
596	GGAGACCG A CGAAGAGC	1360	GCTCTTCG GGCTAGCTACAACGA CGGTCTCC	3735
640	UUGUGGAG A UGGUGGAC	1361	GTCCACCA GGCTAGCTACAACGA CTCCACAA	3736
647	GAUGGUGG A CAACCUGA	1362	TCAGGTG GGCTAGCTACAACGA CCACCATC	3737
650	GGUGGACA A CCUGAGGG	1363	CCCTCAGG GGCTAGCTACAACGA TGTCCACC	3738
688	ACGUGGAG A UGACCGUG	1364	CACGGTCA GGCTAGCTACAACGA CTCCACGT	3739
691	UGGAGAUG A CCGUGGGC	1365	GCCCACGG GGCTAGCTACAACGA CATCTCCA	3740
712	CCCCGAG A CGCUCAAC	1366	GTTGAGCG GGCTAGCTACAACGA CTGCGGGG	3741
719	GACGCUCA A CAUCCUGG	1367	CCAGGATG GGCTAGCTACAACGA TGAGCGTC	3742
731	CCUGGUGG A UACAGGCA	1368	TGCCTGTA GGCTAGCTACAACGA CCACCAGG	3743
746	CAGCAGUA A CUUUGCAG	1369	CTGCAAAG GGCTAGCTACAACGA TACTGCTG	3744
821	AUACCGGG A CCUCCGGA	1370	TCCGGAGG GGCTAGCTACAACGA CCCGGTAT	3745
884	GGCACCG A CCUGGUAA	1371	TTACCAGG GGCTAGCTACAACGA CGGTGCC	3746
911	UGGCCCCA A CGUCACUG	1372	CAGTGACG GGCTAGCTACAACGA TGGGGCCA	3747
929	GCGUGCCA A CAUUGCUG	1373	CAGCAATG GGCTAGCTACAACGA TGGCACGC	3748
948	AUCACUGA A UCAGACAA	1374	TTGTCTGA GGCTAGCTACAACGA TCAGTGAT	3749
953	UGAAUCAG A CAAGUUCU	1375	AGAACTTG GGCTAGCTACAACGA CTGATTCA	3750
968	CUUCAUCA A CGGCUCCA	1376	TGGAGCCG GGCTAGCTACAACGA TGATGAAG	3751
977	CGGCUCCA A CUGGGAAG	1377	CTTCCCAG GGCTAGCTACAACGA TGGAGCCG	3752
1012	AUGCUGAG A UUGCCAGG	1378	CCTGGCAA GGCTAGCTACAACGA CTCAGCAT	3753
1025	CAGGCCUG A CGACUCCC	1379	GGGAGTCG GGCTAGCTACAACGA CAGGCCTG	3754
1028	GCCUGACG A CUCCUGG	1380	CCAGGGAG GGCTAGCTACAACGA CGTCAGGC	3755
1049	UUUCUUUG A CUCUCUGG	1381	CCAGAGAG GGCTAGCTACAACGA CAAAGAAA	3756
1066	UAAAGCAG A CCCACGUU	1382	AACGTGGG GGCTAGCTACAACGA CTGCTTTA	3757
1079	CGUUCCCA A CCUCUUCU	1383	AGAAGAGG GGCTAGCTACAACGA TGGGAACG	3758
1121	CCCCUCA A CCAGUCUG	1384	CAGACTGG GGCTAGCTACAACGA TGAGGGGG	3759
1159	GGAGCAUG A UCAUUGGA	1385	TCCAATGA GGCTAGCTACAACGA CATGCTCC	3760
1175	AGGUAUUG A CCACUCGC	1386	GCGAGTGG GGCTAGCTACAACGA CGATACCT	3761
1240	AUGAGGUG A UCAUUGUG	1387	CACAATGA GGCTAGCTACAACGA CACCTCAT	3762
1258	GGGUGGAG A UCAAUGGA	1388	TCCATTGA GGCTAGCTACAACGA CTCCACCC	3763
1262	GGAGAUCA A UGGACAGG	1389	CCTGTCCA GGCTAGCTACAACGA TGATCTCC	3764

Table 22

1266	AUCAAUGG A CAGGAUCU	1390	AGATCCTG GGCTAGCTACAACGA CCATTGAT	3765
1271	UGGACAGG A UCUGAAAA	1391	TTTTCAGA GGCTAGCTACAACGA CCTGTCCA	3766
1279	AUCUGAAA A UGGACUGC	1392	GCAGTCCA GGCTAGCTACAACGA TTTCAGAT	3767
1283	GAAAAUGG A CUGCAAGG	1393	CCTTGCGG GGCTAGCTACAACGA CCATTTTC	3768
1298	GGAGUACA A CUAUGACA	1394	TGTCATAG GGCTAGCTACAACGA TGTACTCC	3769
1304	CAACUAUG A CAAGAGCA	1395	TGCTCTTG GGCTAGCTACAACGA CATAGTTG	3770
1319	CAUUGUGG A CAGUGGCA	1396	TGCCACTG GGCTAGCTACAACGA CCACAATG	3771
1334	CACCACCA A CCUUCGUU	1397	AACGAAGG GGCTAGCTACAACGA TGGTGGTG	3772
1374	GCAGUCAA A UCCAUCAA	1398	TTGATGGA GGCTAGCTACAACGA TTGACTGC	3773
1412	GUUCCUG A UGUUUUCU	1399	AGAAACCA GGCTAGCTACAACGA CAGGGAAC	3774
1469	CCCUUGGA A CAUUUUC	1400	GGAAATG GGCTAGCTACAACGA TCCAAGGG	3775
1498	UCUACCUA A UGGGUGAG	1401	CTCACCCA GGCTAGCTACAACGA TAGGTAGA	3776
1514	GGUUACCA A CCAGUCCU	1402	AGGACTGG GGCTAGCTACAACGA TGGTAACC	3777
1548	CCGCAGCA A UACUGCG	1403	CGCAGGTA GGCTAGCTACAACGA TGCTGCGG	3778
1568	AGUGGAAG A UGUGCCA	1404	TGGCCACA GGCTAGCTACAACGA CTTCCACT	3779
1586	GUCCCAAG A CGACUGUU	1405	AACAGTCG GGCTAGCTACAACGA CTTGGGAC	3780
1589	CCAAGACG A CUGUUACA	1406	TGTAACAG GGCTAGCTACAACGA CGTCTTGG	3781
1673	UGUCUUUG A UCGGGCCC	1407	GGGCCCGA GGCTAGCTACAACGA CAAAGACA	3782
1686	GCCCCAAA A CGAAUUGG	1408	CCAATTCG GGCTAGCTACAACGA TTTCGGGC	3783
1690	GAAAACGA A UUGGCUUU	1409	AAAGCCAA GGCTAGCTACAACGA TCGTTTTT	3784
1724	UGUGCACG A UGAGUUA	1410	TGAACTCA GGCTAGCTACAACGA CGTGACA	3785
1735	AGUUCAGG A CGGCAGCG	1411	CGCTGCCG GGCTAGCTACAACGA CCTGAAC	3786
1769	CACCUUGG A CAUGGAAG	1412	CTTCCATG GGCTAGCTACAACGA CCAAGGTG	3787
1778	CAUGGAAG A CUGUGGCU	1413	AGCCACAG GGCTAGCTACAACGA CTTCCATG	3788
1790	UGGCUACA A CAUCCAC	1414	GTGGAATG GGCTAGCTACAACGA TGAGCCA	3789
1801	UUCACAG A CAGAUGAG	1415	CTCATCTG GGCTAGCTACAACGA CTGTGGAA	3790
1805	ACAGACAG A UGAGUCAA	1416	TTGACTCA GGCTAGCTACAACGA CTGTCTGT	3791
1813	AGAGUCA A CCCUAUG	1417	CATGAGGG GGCTAGCTACAACGA TGACTCAT	3792
1822	CCCUC AUG A CCAUAGCC	1418	GGCTATGG GGCTAGCTACAACGA CATGAGGG	3793
1925	GCAGCAUG A UGACUUUG	1419	CAAAGTCA GGCTAGCTACAACGA CATGCTGC	3794
1928	GCAUGAUG A CUUUGCUG	1420	CAGCAAAG GGCTAGCTACAACGA CATCATGC	3795
1937	CUUUGCUG A UGACAUCU	1421	AGATGTCA GGCTAGCTACAACGA CAGCAAAG	3796
1940	UGCUGAUG A CAUCUCCC	1422	GGGAGATG GGCTAGCTACAACGA CATCAGCA	3797
1979	GGCAGAAG A UAGAGAUU	1423	AATCTCTA GGCTAGCTACAACGA CTTCTGCC	3798
1985	AGAUAGAG A UUCCCCUG	1424	CAGGGGAA GGCTAGCTACAACGA CTCTATCT	3799
1995	UCCCCUGG A CCACACCU	1425	AGGTGTGG GGCTAGCTACAACGA CCAGGGGA	3800
2033	AGUAGGAG A CACAGAUG	1426	CATCTGTG GGCTAGCTACAACGA CTCCTACT	3801
2039	AGACACAG A UGGACCU	1427	AGGTGCCA GGCTAGCTACAACGA CTGTGTCT	3802
2067	ACCUCAGG A CCCUCCCC	1428	GGGGAGGG GGCTAGCTACAACGA CCTGAGGT	3803
2085	CCCACCAA A UGCCUCUG	1429	CAGAGGCA GGCTAGCTACAACGA TTGGTGGG	3804
2099	CUGCCUUG A UGGAGAAG	1430	CTTCTCCA GGCTAGCTACAACGA CAAGGCAG	3805
2136	UUCAGGG A CUGUACCU	1431	AGGTACAG GGCTAGCTACAACGA CCCTGGAA	3806
2152	UGUAGGAA A CAGAAAAG	1432	CTTTTCTG GGCTAGCTACAACGA TTCCTACA	3807
2189	UGGCGGGA A UACUCUUG	1433	CAAGAGTA GGCTAGCTACAACGA TCCCGCCA	3808
2208	CACCUCAA A UUUAAGUC	1434	GACTTAAA GGCTAGCTACAACGA TTGAGGTG	3809
2222	GUCGGGAA A UUCUGCUG	1435	CAGCAGAA GGCTAGCTACAACGA TTCCCGAC	3810
2237	UGCUGAA A CUUCAGCC	1436	GGCTGAAG GGCTAGCTACAACGA TTCAAGCA	3811
2250	AGCCUGA A CCUUGUC	1437	GACAAAGG GGCTAGCTACAACGA TCAGGGCT	3812
2273	UCCUUUAA A UUCUCCAA	1438	TTGGAGAA GGCTAGCTACAACGA TTAAAGGA	3813
2281	AUUCUCCA A CCCAAAGU	1439	ACTTTGGG GGCTAGCTACAACGA TGAGAAT	3814
2376	GAGAAGAG A CCAAGCUU	1440	AAGCTTGG GGCTAGCTACAACGA CTCTTCTC	3815

Table 22

2417	AGGAGAGG A UGCACAGU	1441	ACTGTGCA GGCTAGCTACAACGA CCTCTCCT	3816
2444	CUUUAGAG A CAGGGACU	1442	AGTCCCTG GGCTAGCTACAACGA CTCTAAAG	3817
2450	AGACAGGG A CUGUAUAA	1443	TTATACAG GGCTAGCTACAACGA CCCTGTCT	3818
2459	CUGUAUAA A CAAGCCUA	1444	TAGGCTTG GGCTAGCTACAACGA TTATACAG	3819
2468	CAAGCCUA A CAUUGGUG	1445	CACCAATG GGCTAGCTACAACGA TAGGCTTG	3820
2482	GUGCAAAG A UUGCCUCU	1446	AGAGGCAA GGCTAGCTACAACGA CTTTGCAC	3821
2494	CCUCUUGA A UAAAAAAAA	1447	TTTTTTAA GGCTAGCTACAACGA TCAAGAGG	3822
2507	AAAAAAAA A CUAGAAAA	1448	TTTCTAG GGCTAGCTACAACGA TTTTTTTT	3823

Input Sequence = AF190725. Cut Site = G/.

Stem Length = 8 . Core Sequence = GGCTAGCTACAACGA

AF190725 (Homo sapiens beta-site APP cleaving enzyme (BACE) mRNA; 2526 bp)

Table 23

Table 23: Human BACE Amberzyme Ribozyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
11	ACGGGUCC G CAGCCGCG	960	GCGGGCUG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGACGCGU	3260
18	CGCAGCCC G CCGGGAG	961	CUCCCGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGGCUUGG	3261
29	CGGGAGCU G CGAGCCGC	962	GCGGCUUG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGCUCCGG	3262
31	GGAGCUGC G AGCCCGCA	963	UCGCGGCU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GCAGCUCC	3263
36	UGGAGCC G CGAGCUGG	964	CCAGCUCG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGCUCGCA	3264
38	CGAGCCGC G AGCUGGAU	965	AUCCAGCU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GCGGCUUG	3265
58	GGUGGCCU G AGCAGCCA	966	UGGCUUGU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGGCCACC	3266
69	CAGCCAAC G CAGCCGCA	967	UGCGGCUU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GUTGGCUG	3267
75	ACGCAGCC G CAGGAGCC	968	GGCUCCUG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGCUGCGU	3268
94	GAGCCCUU G CCCUGGCC	969	GGCAGGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AAGGGCUC	3269
100	UUGCCCCU G CCGCGGCC	970	GCGCGGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGGGGCAA	3270
104	CCUUGCCC G CGCGCGG	971	CGGCGGCG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGGCAGGG	3271
106	CUGCCCGC G CCGCGGCC	972	GCGGCGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GCGGGCAG	3272
109	CCCGGGCC G CCGCGCGC	973	GCGGCGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GCGGCGGG	3273
112	GCGCGGCC G CCGCGCGG	974	CCGCGCGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GCGGCGGC	3274
116	CGCGGCC G CCGGGGGG	975	CCCCCGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGGCGGCG	3275
137	GGGAAGCC G CCACCGGC	976	GCGGUGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGCUUCGC	3276
148	ACGGGCC G CCAUGGCC	977	GGCAUUGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGGCCGCU	3277
153	CCGCGCAU G CCGCGCCC	978	GGGCGCGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AUGGCGGG	3278
157	CCAUGCCC G CCGCUGCC	979	GGGAGGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGGCAUGG	3279
172	CCAGCCCC G CCGGGAGC	980	GCUCGCGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGGGCUUG	3280
183	GGGAGCCC G CGCCCGCU	981	AGCGGGCG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GCGGCUCC	3281
185	GAGCCCGC G CCGCUGGC	982	GCAGCGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GCGGGCUC	3282
189	CCGCGCCC G CUGCCGAG	983	CUGGGCAG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGGCGCGG	3283
192	CGCCCGCU G CCGAGGCU	984	AGCUCUGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGCGGCGG	3284
205	GGCUGGCC G CCGCGGUG	985	CACGCGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGGCAGCC	3285
208	UGGCGGCC G CCGUGCCG	986	CGGACCGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGGGCGCA	3286
213	GCCGCCGU G CCGAUGUA	987	UACAUCGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG ACGGCGGC	3287
216	GCCGUGCC G AUGUAGCG	988	CGCUACAU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG GGCACGCG	3288
250	UCUCCCCU G CUCCCGUG	989	CACGGGAG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGGGGAGA	3289

Table 23

258	GCUCCTCG G CUCUGCGG	990	CCGCAGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACGGGAGC	3290
263	CGUCUCU G CGGAUCUC	991	GAGAUCCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGAGCACG	3291
276	UCUCCTCG G ACCGCUUC	992	AGAGCGGU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGGGGAGA	3292
280	CCUGACC G CUCUCCAC	993	GUGGAGAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGUACAGG	3293
320	AGGGCTCG G CAGGCCCU	994	AGGGCTCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGGGCCU	3294
337	GGCUGUCU G AUGCCCCC	995	GGGGCAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGGACGCC	3295
340	GUCCUGAU G CCCCCAAG	996	CUUGGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUCAGGAC	3296
360	CCUCUCCU G AGAAGCCA	997	UGGCUUCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGGAGAGG	3297
397	GGGCAGGC G CCAGGAC	998	GUCCCTCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GCCUGCCC	3298
420	GGGCCAGU G CGAGCCCA	999	UGGGCTCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACUGGCC	3299
422	GCCAGUGC G AGCCACAG	1000	UCUGGGCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GCACUGGC	3300
437	GAGGCCG G AAGGCCGG	1001	CCGGCCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGGCCUCC	3301
468	CAAGCCU G CCCUGGCU	1002	AGCCAGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGGGCUUG	3302
480	UGGCUCCU G CUGUGGAU	1003	AUCCACAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGGAGCCA	3303
493	GGAUGGC G CGGGAGUG	1004	CACUCCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GCCCAUCC	3304
501	GCGGAGU G CUGCCUGC	1005	GCAGGCAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACUCCCGC	3305
504	GGAGUGCU G CCUGCCCA	1006	UGGGCAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGCACUCC	3306
508	UGCUGCU G CCCACGCG	1007	GCCUGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGGCAGCA	3307
537	AUCCGGCU G CCCCUGCG	1008	CGCAGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGCCGGAU	3308
543	CUGCCCTU G CGCAGCGG	1009	CCGUCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGGGCAG	3309
545	GCCCCUGC G CAGCGGCC	1010	GGCCGCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GCAGGGC	3310
562	UGGGGGC G CCCCCTUG	1011	CAGGGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GCCCCCCA	3311
576	CUGGGCU G CGGUGCC	1012	GGCAGCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGCCCCAG	3312
582	CUGCGCU G CCCCAGGA	1013	UCCGGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGCCGCG	3313
595	GGGAGACC G ACGAAGAG	1014	CUCUUCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGUUCUCC	3314
598	AGACCGAC G AAGAGCCC	1015	GGGCUUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GUCGGUCU	3315
607	AAGAGCC G AGGAGCCC	1016	GGGCUUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGGCUUCU	3316
654	GACAAACU G AGGGGCA	1017	UUGCCCTU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGGUUGUC	3317
690	GUGGAGAU G ACCGUGGG	1018	CCACCGU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUCUCCAC	3318
708	AGCCCCC G CAGACGCU	1019	AGGUCUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGGGGCU	3319
714	CCGCAGAC G CUCACAU	1020	AUGUUGAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GUCUGCG	3320
751	GUAACTUU G CAGUGGGU	1021	ACCCACUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAAGUUC	3321
760	CAGUGGU G CUGCCCCC	1022	GGGGCAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACCCACUG	3322
763	UGGGUGCU G CCCCCCAC	1023	GUGGGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGCACCCA	3323

Table 23

780	CCUUCU G CAUCGUA	1024	UAGCGAUG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AGGAAGGG	3324
785	CUUGCAUC G CUACUACC	1025	GGUAGUAG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG GAUGCAGG	3325
843	GUGUAUGU G CCUACAC	1026	GUGUAGGG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG ACAUACAC	3326
883	UGGGCACC G ACCUGGUA	1027	UACCAGGU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG GGUGCCCA	3327
921	GUCACUGU G CGUGCCAA	1028	UUGGCACG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG ACAGUGAC	3328
925	CUUGUGGU G CCAACAUU	1029	AAUGUUGG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG ACGCACAG	3329
934	CCAACAUU G CUGCCAUC	1030	GAUGGCAG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AAUGUUGG	3330
937	ACAUUGCU G CCAUCACU	1031	AGUGAUGG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AGCAAUGU	3331
946	CCAUCACU G AUCAGAC	1032	GUCUGAUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AGUGAUGG	3332
1006	UGGCCUUAU G CUGAGAUU	1033	AAUCUCAG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AUAGGCCA	3333
1009	CCUAUGCU G AGAUTUGCC	1034	GGCAUCU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AGCAUAGG	3334
1015	CUGAGAUU G CCAGGCCU	1035	AGGCCUUG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AAUCUCAG	3335
1024	CCAGGCCU G ACGACUCC	1036	GGAGUCGU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AGGCCUUGG	3336
1027	GGCCUGAC G ACUCUUG	1037	CAGGGAGU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG GUCAGGCC	3337
1048	CUUUCUUU G ACUCUCUG	1038	CAGAGAGU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AAAGAAG	3338
1092	UUCUCCCU G CAGCUUUG	1039	CAAGCUG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AGGGAGAA	3339
1105	UUUGUGGU G CUGGCUUC	1040	GAAGCCAG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG ACCACAAA	3340
1129	ACCAGUCU G AAGUGCUG	1041	CAGCACUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AGACUGGU	3341
1134	UCUGAAGU G CUGGCCUC	1042	GAGGCCAG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG ACUUCAGA	3342
1158	GGGAGCAU G AUCAUUGG	1043	CCAAUGAU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AUGCUCCC	3343
1174	GAGGUUUC G ACCACUCG	1044	CGAGUGGU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG GAUACCUU	3344
1182	GACCACUC G CUGUACAC	1045	GUGUACAG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG GAGUGGUC	3345
1234	GGUAUUUU G AGGUGAUC	1046	GAUCACCU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AUAUAUCC	3346
1239	UAUGAGGU G AUCAUUGU	1047	ACAAUGAU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG ACCUCAUA	3347
1248	AUCAUUGU G CGGUGUGA	1048	UCCACCCG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG ACAUAGAU	3348
1275	CAGGAUCU G AAAUGGA	1049	UCCAUUUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AGAUCCUG	3349
1286	AAUGGACU G CAAGGAGU	1050	ACUCCUUG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AGUCCAUU	3350
1303	ACAACUUAU G ACAAGAGC	1051	GCUCUUGU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AUAUUGU	3351
1344	CUUGGUUU G CCCAGAA	1052	UUUUGGGG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AAACGAAG	3352
1360	AAGUGUUU G AGCUGCA	1053	UGCAGCUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AAACACUU	3353
1366	UUAGAGCU G CAGUCAAA	1054	UUUGACUG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AGCUUCAU	3354
1411	AGUUCUUU G AUGGUUUC	1055	GAAACCAU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG AGGGAACU	3355
1442	GUUGGUGU G CUGGCAAG	1056	CUUGCCAG GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG ACACCAAGC	3356
1504	UUAUGGGU G AGGUUACC	1057	GGUAACCU GGAGAAACUCC CU UCAAGGACAUUCGUCCGGG ACCCAUUA	3357

Table 23

1526	GUCCUUC G CAUCACCA	1058	UGGUGAUG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GGAAGGAC	3358
1542	AUCCUUC G CAGCAUA	1059	UAUUGCUG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GGAAGGAU	3359
1554	CAAUACCU G CGGCCAGU	1060	ACUGGCCG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGGUUAUG	3360
1588	CCCAAGAC G ACUGUAC	1061	GUAAACGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GUCUUGGG	3361
1603	ACAAGUUT G CCAUCUA	1062	UGAGAUGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AAACUUGU	3362
1672	UUGUCUUT G AUCGGCC	1063	GGCCCGAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AAAGACAA	3363
1682	UCGGGCC G AAAACGAA	1064	UUGGUUUU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GGGCCCGA	3364
1688	CCGAAAAC G AAUUGGCU	1065	AGCCAAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GUUUUGG	3365
1699	UUGGCUUT G CUGUCAGC	1066	GUUGACAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AAAGCCAA	3366
1708	CUGUCAGC G CUUGCCAU	1067	AUGGCAAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GCUGACAG	3367
1712	CAGGCGCU G CCAUGUGC	1068	GCAUCUG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AAGCGCUG	3368
1719	UGCCAUGU G CACGAUGA	1069	UCAUCUG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG ACAUGGCA	3369
1723	AUGUGCAC G AUGAGUUC	1070	GAACUCAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GUGCACAU	3370
1726	UGCAGGAU G AGUUCAGG	1071	CCUGAACU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUCGUGCU	3371
1807	AGACAGAU G AGUCAACC	1072	GGUUGACU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUGAGGCU	3372
1821	ACCCUCAU G ACCAUAGC	1073	GCUAUGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUGAGGCU	3373
1843	UCAUGGCU G CCAUCUGC	1074	GCAGAUGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGCCAUGA	3374
1850	UGCCAUCU G CGCCUCU	1075	AGAGGGCG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGAUGGCA	3375
1852	CCAUCUGC G CCCUCUUC	1076	GAAGAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GCAGAUGG	3376
1863	CUCUUCAU G CUGCCACU	1077	AGUGGCG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUGAAGAG	3377
1866	UUCUUGCU G CCACUCUG	1078	CAGAGUGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGCAUGAA	3378
1874	GCCACUCU G CCUAUGG	1079	CCAUGAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGAGUGGC	3379
1895	UCAGUGGC G CUGCCUCC	1080	GGAGGCAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GCCACUGA	3380
1898	GUGGGGCU G CCUCGCU	1081	AGCGGAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGCGCCAC	3381
1904	CUGCCUCC G CUGCCUCC	1082	GCAGGCAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GGAGGCAG	3382
1907	CCUCGCGU G CCUGCGCC	1083	GGCGCAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGCGGAGG	3383
1911	CGCUGCCU G CGCCAGCA	1084	UGCUGGCG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGGCAGCG	3384
1913	CUGCCUCC G CCAGCAGC	1085	GCUCUGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GCAGGCAG	3385
1924	AGCAGCAU G AUGACUUT	1086	AAAGUCAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUGCUGCU	3386
1927	AGCAUGAU G ACUUGCU	1087	AGCAAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUCAUGCU	3387
1933	AUGACUUT G CUGAUGAC	1088	GUCAUCAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AAAGUCAU	3388
1936	ACUUGCU G AUGACAUC	1089	GAUGUCAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGCAAGAU	3389
1939	UUGCUGAU G ACAUCUCC	1090	GGAGAUGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUCAGCAA	3390
1950	AUCUCCCU G CUGAAGUG	1091	CACUUCAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGGGAGAU	3391

Table 23

1953	UCCUUGCU G AAGUGAGG	1092	CCUCACUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGCAGGGA	3392
1958	GUUGAAGU G AGGAGGCC	1093	GGCCUCCU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACUUCAGC	3393
2087	CACCAAU G CCUCUGCC	1094	GGCAGAGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUUUGGUG	3394
2093	AUGCCUCU G CCUUGAUG	1095	CAUCAAGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGAGGCAU	3395
2098	UCUGCCUU G AUGGAGAA	1096	UUCUCCAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAGGCAGA	3396
2179	AGCACUCU G CUGGCGGG	1097	CCGCGCAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGAGUGCU	3397
2227	GAUAUUCU G CUGCUUGA	1098	UCAAGCAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGAUUUC	3398
2230	AUUCUGCU G CUGAAAC	1099	GUUCAAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGCAGAAU	3399
2234	UGCUGCUU G AAACUUCA	1100	UGAAGUUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAGCAGCA	3400
2248	UCAGCCCU G AACCUUUG	1101	CAAAGUUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGGGCUGA	3401
2329	CAUCACAC G CAGGUAAC	1102	GUAAACUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG GUGUGAUG	3402
2393	GUUCCCU G CUGGCCAA	1103	UUGGCGAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGGGAAC	3403
2419	GAGAGGAU G CACAGUUU	1104	AAACUGUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUCCUUC	3404
2428	CACAGUUU G CUUUUGC	1105	GCAAUUAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAACUGUG	3405
2435	UGCUAUUU G CUUUAGAG	1106	CUCUAAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAUAGCA	3406
2476	ACAUUGGU G CAAAGAUU	1107	AAUCUUUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACCAAUGU	3407
2485	CAAAGAUU G CCUCUUGA	1108	UCAAGAGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAUCUUUG	3408
2492	UGCCUCUU G AAUUA AAA	1109	UUUUAAUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAGAGGCA	3409
219	GUGCCGAU G UAGCGGGC	1110	GCCCGCUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUCGGCAC	3410
483	CUCCUGCU G UGGAUGGG	1111	CCCAUCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGCAGGAG	3411
634	GCAGCUUU G UGGAGAU	1112	CAUCUCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAAGCUGC	3412
804	AGGAGCUU G UCCAGCAC	1113	GUGCUGGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGTUGCCU	3413
835	GGAGGGU G UGUUUGUG	1114	CACAUACA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACCCUUCC	3414
837	AAGGUGU G UAUGUGCC	1115	GGCACAUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACACCCUU	3415
841	GUGUGUAU G UGCCCUAC	1116	GUAGGGCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUACACAC	3416
919	ACGUCACU G UGCGUGCC	1117	GGCAGCGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGUGACGU	3417
1100	GCAGCUUU G UGGUGCUG	1118	CAGCACCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAAGCUGC	3418
1144	UGGCCUCU G UCGGAGGG	1119	CCUCCCGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGAGGCCA	3419
1185	CACUGCUU G UACACAGG	1120	CCUGUGUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGCGAGUG	3420
1246	UGAUCAUU G UGCGGGUG	1121	CACCCGCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAUGAUCA	3421
1315	AGAGCAUU G UGGACAGU	1122	ACUGUCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAUGCUCU	3422
1356	AAGAAAGU G UUUGAAGC	1123	GCUUCAAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACUUCUUU	3423
1440	CAGCUGGU G UGCUGGCA	1124	UGCCAGCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACCAGCUG	3424
1570	UGGAAGAU G UGGCCACG	1125	CGUGGCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUCUUCCA	3425

Table 23

1592	AGACGACU G UUAACAAGU	1126	ACUUGUAA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGUCGUCU	3426
1630	CGGGCACU G UUAUGGGA	1127	UCCCAUAA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGUGCCCG	3427
1642	UGGGAGCU G UUAUCAUG	1128	CAUGAUAA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGCUCCCA	3428
1666	UCUACGUU G UCUIUGAU	1129	AUCAAGA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AACGUAGA	3429
1702	GCUUUGCU G UCAGCGCU	1130	AGCGCUGA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGCAAAGC	3430
1717	CUUGCCAU G UGCACGAU	1131	AUCGUGCA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AUGGCAAG	3431
1759	GCCUUIUJ G UCACCTUG	1132	CAAGGUGA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AAAAGGGC	3432
1781	GGAAGACU G UGGCUACA	1133	UGUAGCCA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGUCUUC	3433
1834	UAGCCUUAU G UCAUGGCU	1134	AGCCAUUA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AUAGGCUA	3434
1884	CUCAUGGU G UGUCAGUG	1135	CACUGACA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG ACCAUGAG	3435
1886	CAUGGUGU G UCAGUGGC	1136	GCCACUGA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG ACACAUUG	3436
2048	UGGCACCU G UGGCCAGA	1137	UCUGGCCA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGGUGCCA	3437
2139	CAGGGACU G UACCUGUA	1138	UACAGGUA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGUCCUG	3438
2145	CUGUACCU G UAGGAAAC	1139	GUUUCUUA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGUACAG	3439
2256	GRACCUUU G UCCACCAU	1140	AUGGUGGA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AAAGSUUC	3440
2346	CUUGGCGU G UGUCCUG	1141	CAGGGACA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG ACGCCAAG	3441
2348	UGGCGUGU G UCCUGUG	1142	CACAGGGA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG ACACGCCA	3442
2354	GUGUCCCU G UGGUACCC	1143	GGGUACCA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGGACAC	3443
2385	CCAAGCUU G UUUCCUG	1144	CAGGAAA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGCUUUG	3444
2453	CAGGGACU G UAUAAACA	1145	UGUUUAUA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGUCCUG	3445
14	CGUCCGCA G CCCGCCG	1146	CGGGCGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UCGGAGCG	3446
26	GCCCGGA G CUGCGAGC	1147	GCUCGAG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UCCCGGGC	3447
33	AGCUGCGA G CCGGAGC	1148	GCUCGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UCGCAGCU	3448
40	AGCCCGGA G CUGGAUUA	1149	UAAUCAG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UCGCGGCU	3449
51	GGAUUAUG G UGGCCUGA	1150	UCAGGCCA GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG CAUAAUCC	3450
54	UUAUGGUG G CCUGAGCA	1151	UGCUCAGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG CACCAUAA	3451
60	UGGCCUGA G CAGCCAAC	1152	GUUGGUG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UCAGGCCA	3452
63	CCUGAGCA G CCAAGCA	1153	UGGUGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGCUACAG	3453
72	CCAACGCA G CCGCAGGA	1154	UCCUGCG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGCGUUGG	3454
81	CGCAGGA G CCCGAGC	1155	GCUCGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UCCUGCGG	3455
88	AGCCCGGA G CCUUGCC	1156	GGCAAGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UCCGGGCU	3456
134	CAGGGAA G CCGCCACC	1157	GGUGCGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UUCUCCUG	3457
144	CGCACCG G CCGCCAU	1158	AUGGCGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG CGUGGCG	3458
167	CCUCCCA G CCCGCCG	1159	CGGCGGG GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGGAGGG	3459

Table 23

179	CGCCGGGA G CCGCGCC	1160	GGCGCGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCCGGGG	3460
198	CUGCCAG G CUGCCGC	1161	GGGCCAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGGGCAG	3461
202	CCAGGUG G CCGCGCC	1162	GGCGCGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGCCUGG	3462
211	CCGCGCC G UGCCGAG	1163	CAUCGGCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GCGGCGG	3463
222	CCGAUGA G CGGGUCC	1164	GGAGCCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UACAUCGG	3464
226	UGUAGCG G CUCGGAU	1165	AUCCGGAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCGCUACA	3465
239	GGAUCCA G CCUCUCC	1166	GGGAGAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGGGAUCC	3466
256	CUGCUCC G UGCUGC	1167	GCAGAGCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGGAGCAG	3467
290	UCUCCACA G CCGGACC	1168	GGUCGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGUGGAGA	3468
304	ACCGGGG G CUGGCCA	1169	UGGCGCAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCCAGGU	3469
308	GGGGCUG G CCCAGGC	1170	GCCUGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGCCCC	3470
315	GGCCAGG G CCUGGCG	1171	CUGCAGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGGGCC	3471
324	CCUGCAG G CCUGGCG	1172	CGCCAGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGAGGG	3472
330	AGGCCUG G CGUCCUGA	1173	UCAGGAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGGGCCU	3473
332	GCCUGGC G UCCUGAG	1174	CAUCAGGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GCCAGGGC	3474
348	GCCCCAA G CUCCUCU	1175	AGAGGGAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUGGGGGC	3475
365	CCUGAGAA G CCACAGC	1176	GUGUGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUCUCAGG	3476
372	AGCACCA G CACCACC	1177	GGUGGUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGGUGGCU	3477
391	ACUUGGG G CAGCGCC	1178	GGCGCUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCCAGU	3478
395	GGGGCAG G CGCCAGG	1179	CCUGGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGCCCC	3479
410	GGACGGAC G UGGGCCAG	1180	CUGGCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GUCCGUCC	3480
414	GGACUGG G CCAGUGC	1181	CGCACUGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCACGUCC	3481
418	GUGGGCA G UGCAGCC	1182	GGCUCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGGCCAC	3482
424	CAGUGCGA G CCAGAGG	1183	CCUCUGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCGCACUG	3483
433	CCAGAGG G CCGRAGG	1184	CCUUGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCUCUGGG	3484
441	GCCGAG G CCGGGCC	1185	GGCCCCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUUCGGGC	3485
447	AGGCCGG G CCGACAU	1186	AUGGUGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCGGGCCU	3486
457	CCACCAUG G CCAAGCC	1187	GSCUUGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAUGGUGG	3487
463	UGGCCAA G CCUGCCC	1188	GGGCAGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUGGGCCA	3488
474	CUGCCUG G CUCCUGU	1189	AGCAGGAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGGGCAG	3489
491	GUGGAUG G CGCGGAG	1190	CUCCCGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCAUCAC	3490
499	GCGGGGA G UGCUGCU	1191	AGCAGCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCCGGC	3491
515	UGCCACG G CACCAGC	1192	GCUGGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGUGGCA	3492
522	GGCACCA G CACGGAU	1193	AUGCCUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGGGUGCC	3493

Table 23

527	CCAGCAG G CAUCCGGC	1194	GCCGAUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGUGCUGG	3494
534	GGCAUCC G CUGCCCCU	1195	AGGGCAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGGAUGCC	3495
548	CCUGCGCA G CGGCTUGG	1196	CCAGCCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCGCAGG	3496
551	GGCAGCG G CCUGGGGG	1197	CCCCAGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGUGCGC	3497
560	CCUGGGG G CGCCCCC	1198	GGGGGCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCCAGG	3498
573	CCCCUGG G CUGGGGU	1199	AGCCGCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCAGGG	3499
579	GGCUGCG G CUGCCCCG	1200	CGGGCAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGCAGCC	3500
603	GACGAAG G CCGAGGA	1201	UCCUCGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCUCGUC	3501
612	CCGAGGA G CCGGCCG	1202	CGGCCGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCUCGG	3502
617	GGAGCCG G CCGAGGG	1203	CCUCCGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGGCUCC	3503
626	CCGAGGG G CAGCUUG	1204	CAAAGCUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCUCCGG	3504
629	GAGGGCA G CUUGUGG	1205	CCACAAAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCCCUCC	3505
643	UGGAGAUG G UGACAAC	1206	GUUGCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAUCUCCA	3506
659	CCUGAGG G CAAGUCGG	1207	CCGACUUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCUCAGG	3507
663	AGGGGCA G UCGGGCA	1208	UGCCCCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCCCCU	3508
669	AAGUCGG G CAGGCUA	1209	UGCCCUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCACUU	3509
674	GGGCGAG G CUACUACG	1210	CGUAGUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUGCCCC	3510
682	GUACUAC G UGGAGUG	1211	CAUCUCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GUAGUAG	3511
694	AGAUGAC G UGGCAGC	1212	GCUGCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCACGGUC	3512
698	GACCGUG G CAGCCCC	1213	GGGGGUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCCACG	3513
701	CGUGGCA G CCCCCCG	1214	GCGGGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCCACG	3514
727	ACAUCCUG G UGAUACA	1215	UGUAUCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGGAUG	3515
737	GAUACAG G CAGCAGUA	1216	UACUGCUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGUAUCC	3516
740	UACAGGCA G CAGUAACU	1217	AGUACUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCCUGUA	3517
743	AGCAGCA G UAACUUUG	1218	CAAAGUUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCUGCU	3518
754	ACUUUGCA G UGGUGCU	1219	AGCACCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCAAAU	3519
758	UGCAGUG G UGUGCCC	1220	GGGACCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCACUGCA	3520
798	UACCAGAG G CAGCUGC	1221	GACAGCUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUCUGGUA	3521
801	CAGAGGCA G CUGUCCAG	1222	CUGGACAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCCUCUG	3522
809	CGUGUCCA G CACAUACC	1223	GGUAUGUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGGACAGC	3523
833	CCGGAAG G UGUGAUG	1224	CAUACACA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCUCCCG	3524
857	CACCCAG G CAAGUGG	1225	CCCACUUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCUGGGUG	3525
861	CAGGGCA G UGGGAAG	1226	CCUCCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUGCCUG	3526
873	GAAGGGA G CUGGGCAC	1227	GUGCCAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCCUUC	3527

Table 23

878	GGAGCUGG G CACCGACC	1228	GGUCGGUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCAGCUCC	3528
889	CCGACCUG G UAAGCAUC	1229	GAUGCUUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGGUCCG	3529
893	CCUGGUAA G CAUCCCC	1230	GGGGGAUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUACCAGG	3530
905	CCCCCAUG G CCCCAACG	1231	CGUUGGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAUGGGGG	3531
913	GGCCCAAC G UCACUGUG	1232	CACAGUGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GUUGGGGC	3532
923	CACUGUGC G UGCCAACA	1233	UGTUGGCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GCACAGUG	3533
957	UCAGACAA G UUCUUAU	1234	AUGAAGAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUGUCUGA	3534
971	CAUCAACG G CUCCAACU	1235	AGUUGGAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGUUGAUG	3535
986	CUGGGAAG G CAUCCUGG	1236	CCAGGAUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUUCCCAG	3536
996	AUCCUGGG G CUGGCCUA	1237	UAGGCCAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCAGGAU	3537
1000	UGGGGCGUG G CCUAUGCU	1238	AGCAUAGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGCCCCA	3538
1020	AUUGCCAG G CCUGACGA	1239	UCGUCAGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGGCAAU	3539
1038	UCCUCUGA G CCUUCUUA	1240	AAGAAAGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCAGGGA	3540
1057	ACUCUCUG G UAAAGCAG	1241	CUGCUUUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGAGAGU	3541
1062	CUGGUAAA G CAGACCCA	1242	UGGGUCUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUUACCAG	3542
1072	AGACCCAC G UUCCCAAC	1243	GUUGGGAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GUGGGUCU	3543
1095	UCCUCUGA G CUUUGUGG	1244	CCACAAAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCAGGGA	3544
1103	GCUUUGUG G UGUGGCGU	1245	AGCCAGCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CACAAAGC	3545
1109	UGGUGCUG G CUUCCCC	1246	GGGGGAAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGCACCA	3546
1125	CUCAACCA G UCUGAAGU	1247	ACUUCAGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGGUUGAG	3547
1132	AGUCUGAA G UGUGGCC	1248	GGCCAGCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUCAGACU	3548
1138	AAGUGCUG G CCUCUGUC	1249	GACAGAGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGCACUU	3549
1154	CGGAGGGA G CAUGAUCA	1250	UGAUCUAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCCUCCG	3550
1169	CAUUGGAG G UAUCGACC	1251	GGUCGAUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUCCAUG	3551
1193	GUACACAG G CAGUCUCU	1252	AGAGACUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGUGUAC	3552
1196	CACAGGCA G UCUCUGGU	1253	ACCAGAGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCCUGUG	3553
1203	AGUCUCUG G UAUACACC	1254	GGUGUAUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGAGACU	3554
1218	CCCAUCCG G CGGGAGUG	1255	CACUCCCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGGAUGGG	3555
1224	CGCGGGGA G UGUUAUA	1256	UAUACCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCCGCCG	3556
1227	CGGGAGUG G UAUUAUGA	1257	UCAUAUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CACUCCCG	3557
1237	AUUAUGAG G UGAUCAUU	1258	AAUGAUCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUCAUAAU	3558
1252	UUGUGCGG G UGGAGAU	1259	GAUCCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCGCACAA	3559
1293	UGCAAGGA G UACAACUA	1260	UAGTUGUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCUUGCA	3560
1310	UGACAAGA G CAUUGUGG	1261	CCACAAUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCUUGUCA	3561

Table 23

1322	UGUGGACA G UGGACCA	1262	UGUGCCA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGUCCACA	3562
1325	GGACAGUG G CACCACCA	1263	UGUGGUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CACUGUCC	3563
1340	CAACCUUC G UUUGCCA	1264	UGGCAAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG GAAGGUUG	3564
1354	CCAAGAAA G UGUUGAA	1265	UUCAAA CA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUUCUUGG	3565
1363	UGUUUGAA G CUGCAGUC	1266	GACUGCAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUCAACA	3566
1369	AAGCUGCA G UCAAAUCC	1267	GGAUUUGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGCAGCUU	3567
1384	CCAUCAAG G CAGCCUCC	1268	GGAGGCUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUUGAUGG	3568
1387	UCAAGGCA G CUCCUCC	1269	GGAGGAGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGCCUUGA	3569
1404	ACGGAGAA G UUCUUGA	1270	UCAGGAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUCUCCGU	3570
1415	CCCUUGAUG G UUUCUGGC	1271	GCCAGAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CACAGGG	3571
1422	GGUUUCUG G CUAGGAGA	1272	UCUCCUAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CAGAAACC	3572
1431	CUAGGAGA G CAGCUGGU	1273	ACCAGCUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGCUCUAG	3573
1434	GGAGAGCA G CUGGUGUG	1274	CACACCAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGCUCUCC	3574
1438	AGCAGCUG G UGUGCUGG	1275	CCAGCACA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CAGCUCU	3575
1446	GUGUGCUG G CAGGCAGG	1276	CCUGCUUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CAGCACAC	3576
1450	GCUGGCAA G CAGGCACC	1277	GGUGCCUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUGCCAGC	3577
1454	GCAAGCAG G CACCACCC	1278	GGUGGUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUGCUUGC	3578
1480	UUUUCCCA G UCAUCUCA	1279	UGAGUGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGGGAAA	3579
1502	CCUAAUGG G UGAGGUUA	1280	UAACCUCA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCAUUAAG	3580
1507	UGGUGAG G UUAACCAAC	1281	GUUGGUA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUCACCCA	3581
1518	ACCAACCA G UCCUCCG	1282	CGAAGGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGGUUGU	3582
1545	CUUCCGCA G CAAUACCU	1283	AGGUUUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGCGBAAG	3583
1557	UACUUGG G CCAGUGGA	1284	UCCACUGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CGCAGGUA	3584
1561	UGCGGCA G UGGAAGAU	1285	AUCUCCA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGGCCGCA	3585
1573	AAGAUUG G CCAUGUCC	1286	GGACUGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CACAUCU	3586
1578	GUGGCCAC G UCCCAAGA	1287	UCUUGGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG GUGGCCAC	3587
1599	UGUUAACA G UUUGCCAU	1288	AUGGCAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUGUAACA	3588
1614	AUCUCACA G UCAUCCAC	1289	GUGGAUGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGUGAGAU	3589
1625	AUCCAGG G CACUGUUA	1290	UAACAGUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCGUGGAU	3590
1639	UUUUGGA G CUGUUAUC	1291	GAUAACAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UCCCAUAA	3591
1655	CAUGGAG G CUUCUACG	1292	CGUAGAAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCUCCAUG	3592
1663	GCUCUAC G UUGUCUUU	1293	AAAGACAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG GUAGAAGC	3593
1678	UUGAUCGG G CCCGAAA	1294	UUUUCGGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCGAUCAA	3594
1694	ACGAUUG G CUUUGCUG	1295	CAGCAAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CAAUUCGU	3595

Table 23

1706	UGCUGUCA G CGCUUGCC	1296	GGCAAGCG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UGACAGCA	3596
1728	CACGAUGA G UUCAGGAC	1297	GUCCUGAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UCAUCGUG	3597
1738	UCAGGACG G CAGCGGUG	1298	CACCGCUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CGUCUGA	3598
1741	GGACGGCA G CGGUGGAA	1299	UCCACCG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UGCCGUCC	3599
1744	CGCAGCG G UGGAAGGC	1300	GCCUCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CGCUGCCG	3600
1751	GGUGGAAG G CCCUUUG	1301	CAAAAGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CUUCACC	3601
1784	AGACUGUG G CUACAACA	1302	UGTUGAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CACAGUCU	3602
1809	ACAGAUGA G UCAACCCU	1303	AGGTUGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UCAUCUGU	3603
1828	UGACCAUA G CCUAUGUC	1304	GACAUAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UAUGUCA	3604
1840	AUGUCAUG G CUGCCAUC	1305	GAUGGAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAUGACA	3605
1882	GCTUCAUG G UGUGUCAG	1306	CUGACACA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAUGAGGC	3606
1890	GUGUGUCA G UGGCGCUG	1307	CAGCGCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UGACACAC	3607
1893	UGUCAGUG G CGCUGCCU	1308	AGGACG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CACUGACA	3608
1917	CUGCGCCA G CAGCAUGA	1309	UCAUCUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UGGCGCAG	3609
1920	CGCCAGCA G CAUGAUGA	1310	UCAUCAUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UGCUGCG	3610
1956	CUGCUGAA G UGAGGAGG	1311	CCUCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UUCAGCAG	3611
1964	GUGAGGAG G CCCAUGGG	1312	CCAUGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CUCCUCAC	3612
1972	GCCCAUGG G CAGAAGAU	1313	AUCUCUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CCAUGGCG	3613
2006	ACACCUCC G UGGUUCAC	1314	GUGAACA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG GGAGGUGU	3614
2009	CCUCCGUG G UUCACUUU	1315	AAAGUAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CACGGAGG	3615
2019	UCACUUUG G UCACAAGU	1316	ACUUGUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAAAGUGA	3616
2026	GGUCACAA G UAGGAGAC	1317	GUCUCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UUGUGACC	3617
2042	CACAGAUG G CACUUGUG	1318	CACAGGUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAUCUGUG	3618
2051	CACUUGUG G CCAGAGCA	1319	UGCUCUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CACAGGUG	3619
2057	UGGCCAGA G CACUUCAG	1320	CUGAGGUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UCUGGCCA	3620
2114	AGGAAAG G CUGGCNAG	1321	CUUGCCAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CUUUUCU	3621
2118	AAAGCUG G CAAGGUGG	1322	CCACUUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAGCCUUU	3622
2123	CUGGCAAG G UGGGUUCC	1323	GGAAACCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CUUGCCAG	3623
2127	CAAGGUGG G UUCAGGG	1324	CCUUGAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CCACCUUG	3624
2172	AGAAAGAA G CACUUGC	1325	GCAGAGUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UUCUUUCU	3625
2183	CUCUGCUG G CGGGAUA	1326	UAUUCGCG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAGCAGAG	3626
2198	UACUCUUG G UCACCUCA	1327	UGAGGUGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAAAGAGUA	3627
2214	AAAUUUA G UCGGGAAA	1328	UUUCCCGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UUAUUUU	3628
2243	AAACUUA G CCUGAAC	1329	GUUCAGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UGAAGUUU	3629

Table 23

2288	AACCCAAA	G	UAUUCUUC	1330	GAAGAAUA	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	UUUGGUU	3630
2305	UUUCUUUA	G	UUUCAGAA	1331	UUUCGAAA	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	UNAGAAAA	3631
2314	UUUCAGAA	G	UACUGGCA	1332	UGCCAGUA	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	UUCUGAAA	3632
2320	AAGUACUG	G	CAUCACAC	1333	GUGUGAUG	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CAGUACUU	3633
2333	ACACGCAG	G	UUACCUUG	1334	CAAGGUAA	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CUGCGUGU	3634
2342	UUACCUUG	G	CGUGUGUC	1335	GACACACG	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CAAGGUAA	3635
2344	ACCUUGGC	G	UGUGUCCC	1336	GGGACACA	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	GCCAAGGU	3636
2357	UCCCUUGG	G	UACCCUGG	1337	CCAGGGUA	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CACAGGGA	3637
2365	GUACCCUG	G	CAGAGAAG	1338	CUUCUCUG	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CAGGUAAC	3638
2381	GAGACCAA	G	CUUGUUUC	1339	GAACAAG	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	UUGGUCUC	3639
2397	CCUUGCUG	G	CCAAAGUC	1340	GACUUUG	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CAGCAGGG	3640
2403	UGGCCAAA	G	UCAGUAGG	1341	CCUACUGA	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	UUUGGCCA	3641
2407	CAAAAGUCA	G	UAGGAGAG	1342	CUCUCCUA	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	UGACUUUG	3642
2424	GAUGCACA	G	UUUGCUAU	1343	AUAGCAA	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	UGUGCAUC	3643
2463	AUAAACAA	G	CCUAACAU	1344	AUGUAGG	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	UUGUUUAU	3644
2474	UAACAUGG	G	UGCAAAGA	1345	UCUUUGCA	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CAUGUUA	3645
22	GCCCGCCC	G	GGAGCUGC	1449	GCAGCUCC	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	GGCGCGGC	3646
23	CCCGCCCG	G	GAGCUGCG	1450	CGCAGCUC	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CGGCGGGG	3647
24	CCGCCCGG	G	AGCUGCGA	1451	UCGACGU	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CGGCGGGG	3648
43	CGCGAGCU	G	GAUUAUGG	1452	CCAUAUC	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	AGCUCGCG	3649
44	GGGAGCUG	G	AUUAUGGU	1453	ACCAUAU	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CAGCUCGC	3650
50	AUUAUGGU	G	GUGGCCUG	1454	CAGGCCAC	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	AUAUCCA	3651
53	AGCCCGCA	G	GCCUGAGC	1455	GCUCAGGC	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	ACCAUAU	3652
78	AGCCCGCA	G	GAGCCCGG	1456	CCGGCUC	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	UGCGGCUG	3653
79	AGCCCGCAG	G	AGCCCGGA	1457	UCGGGCU	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CUGCGGCU	3654
85	AGGAGCCC	G	GAGCCCUU	1458	AAGGGCUC	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	GGGCUCCU	3655
86	GGAGCCCG	G	AGCCCUUG	1459	CAAGGGCU	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CGGGCUCC	3656
119	GGCCCGCC	G	GGGGGACC	1460	GGUCCCC	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	GGCGGGCG	3657
120	GGCCCGCC	G	GGGGACCA	1461	UGGUCCCC	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CGGCGGGC	3658
121	CCCGCCGG	G	GGGACCAG	1462	CUGGUCCC	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CCTGGCGG	3659
122	CGCCCGGG	G	GGACCAGG	1463	CCUGGUCC	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CCTGGCGG	3660
123	CGCCCGGG	G	GACCAGGG	1464	CCCTUGGU	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CCTGGCGG	3661
124	GCCGGGGG	G	ACCAGGGA	1465	UCCCTUGU	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	CCTGGCGG	3662
129	GGGACCA	G	GGAAGCCG	1466	CGGCUCC	GGAGGAAACUCC	CU	UCAAGGACAUCGUCGCGG	UGGUCCCC	3663

Table 23

130	GGGACCAG G GAAGCCGC	1467	GCGGCTUC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CUGGUCCC	3664
131	GGACCHAG G AAGCCGCC	1468	GGCGGCTU GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CCUGGUCC	3665
143	CCGCCACC G GCCCGCCA	1469	UGGCGGGC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG GGUUGCGG	3666
175	GCCCCGCC G GGAGCCCG	1470	CGGGCUCC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG GCGCGGGC	3667
176	CCCCGCCG G GAGCCCGC	1471	GCGGGCUC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CCGCGGGG	3668
177	CCCCCGCG G AGCCCGCG	1472	CGCGGGCU GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CCGCGGGG	3669
197	GCUGCCCA G GCUGGCCG	1473	CGCCGAGC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG UGGGCAGC	3670
201	CCCAGGCU G GCCGCCGC	1474	GCGCGGGC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG AGCCUGGG	3671
224	GAUGUAGC G GCUCCCGG	1475	CCGAGGCC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG GCUACAU	3672
225	AUGUAGCG G GCUCCGGA	1476	UCCGGAGC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CGCUACAU	3673
231	CGGGCUCC G GAUCCCGA	1477	CUGGAUC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG GGAGCCCG	3674
232	GGGCUCCG G AUCCCGAG	1478	CUUGGAU GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CGAGAGCC	3675
265	UGCUCUGC G GAUCUCCC	1479	GGGGAUC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG GCAAGACA	3676
266	GCUCUGCG G AUCUCCCC	1480	GGGAGAU GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CGCAGAGC	3677
294	CACAGCCC G GACCCGGG	1481	CCCGGUUC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG GGGCUGUG	3678
295	ACAGCCCG G ACCCGGGG	1482	CCCCGGU GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CCGGGCUGU	3679
300	CCGAGCCC G GGGGCUUG	1483	CCAGCCC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG GGGUCCCG	3680
301	CGGACCCG G GGGCUGGC	1484	GCCAGCCC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CCGGUCCG	3681
302	GGACCCGG G GGCUGGCC	1485	GGCCAGCC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CCGGGUCC	3682
303	GACCCGGG G GCUGGCCC	1486	GGGCCAGC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CCGGGGUC	3683
307	CGGGGGCU G GCCCAGGG	1487	CCCUGGC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG AGCCCCCG	3684
313	CUGGCCCA G GGCCUUGC	1488	GCAGGGCC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG UGGGCCAG	3685
314	UGGCCCGA G GCCCUGCA	1489	UGCAGGGC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CUGGGCCA	3686
323	GCCCUGCA G GCCCUGGC	1490	GCCAGGGC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG UGCAGGGC	3687
329	CAGGCCCCU G GCGUCCUG	1491	CAGGAGCC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG AGGCGCUG	3688
362	UCUCCUGA G AAGCCACC	1492	GGUGGCTU GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG UGCAGAGA	3689
382	ACCACCCA G ACUUGGGG	1493	CCCCAAGU GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG UGGGUGGU	3690
387	CCAGACTU G GGGGCAGG	1494	CCUGCCCC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG AAGUCUGG	3691
388	CAGACTUG G GGCAGGCC	1495	GCCUGCCC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CAAGUCUG	3692
389	AGACUUGG G GGCAGGCG	1496	CGCUGCC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CCAAGUCU	3693
390	GACUUGGG G GCAGGCGC	1497	GCGCCUGC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CCAAGUCU	3694
394	UGGGGGCA G GCGCCAGG	1498	CCUGGCGC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CCGGAGUC	3695
401	AGCGGCCA G GGACGGAC	1499	GUCCGUCC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG UGGGCGCU	3696
402	GGCGCCAG G GACGGACG	1500	CGUCCGUC GGAGGAAACUCC CU UCAAGGACAUUCGUCGGG CUGGCGCC	3697

Table 23

403	GGCCAGG G ACGACGU	1501	ACUCCGU GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CCUGGCGC	3698
406	CCAGGGAC G GACGUGG	1502	CCCACUC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG GUCCUUG	3699
407	CAGGGACG G ACGUGGC	1503	GCCACGU GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CGUCCUG	3700
412	ACGGACGU G GGCAGUG	1504	CACUGCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG ACGUCCU	3701
413	CGGACGUG G GCCAGUG	1505	GCACUGC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CACGUCCG	3702
429	CGAGCCCA G AGGCCCG	1506	CGGCCCU GGAGAAAUCC CU UCAAGGACAUCGUCCGGG UGGGCUUG	3703
431	AGCCAGA G GGCCCGAA	1507	UUGGGCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG UCUGGCGU	3704
432	GCCAGAG G GCCGAAG	1508	CUUCGGC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CUCUGGGC	3705
440	GGCCCGAA G GCGGGGC	1509	GCCCGGC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG UUCGGGCC	3706
444	CGAAGGC G GGCCCCAC	1510	GUGGGCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG GGCCUUCG	3707
445	GAAGGCC G GGCCACC	1511	GGUGGCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CGGCCUUC	3708
446	AAGGCCG G GCCACCA	1512	UGUGGCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CCGGCCUU	3709
456	CCACCAU G GCCAAGC	1513	GCUUGGC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG AUGGUGG	3710
473	CCUGCCCU G GCUCUGC	1514	GCAGAGC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG AGGCAGG	3711
485	CCUGCUGU G GAUGGCG	1515	CGCCAU C GGAGAAAUCC CU UCAAGGACAUCGUCCGGG ACAGAGG	3712
486	CUGCUGUG G AUGGCGC	1516	GCGCCAU GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CACAGCAG	3713
489	CUGUGGAU G GCGCGGG	1517	CCCGGCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG AUCCACAG	3714
490	UGUGGAUG G GCGGGGA	1518	UCCCGCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CAUCCACA	3715
495	AUGGGCG G GAGUGCU	1519	AGCACUC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG GCGCCAU	3716
496	UGGGCGG G GAGUGCUG	1520	CAGCACU GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CGCGCCA	3717
497	GGCGCGG G AGUGCUG	1521	GCAGCAU GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CCGGCC	3718
514	CUGCCAC G GCACCCAG	1522	CUGGUGC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG GUGGCAG	3719
526	CCAGCAC G GCAUCCGG	1523	CCGGAUG GGAGAAAUCC CU UCAAGGACAUCGUCCGGG GUGCUGG	3720
533	CGGCAUC G GCUCCCG	1524	GGGCAGC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG GGAUGCCG	3721
550	UGCGCAGC G GCUUGGG	1525	CCCGGCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG GCUUGCA	3722
555	AGCGGCCU G GGGGGCG	1526	GCGCCCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG AGGCGCU	3723
556	GCGGCCUG G GGGGCGC	1527	GGGCCCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CAGGCGC	3724
557	CGGCTUGG G GGGGCGC	1528	GGGCCCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CCAGGCG	3725
558	GGCTUGG G GGGGCGC	1529	GGGCGCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CCCAGGC	3726
559	GCCUGGG G GGGGCGC	1530	GGGGGCG GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CCCCAGC	3727
570	GCCCCCU G GGGCUGG	1531	CGCAGCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG AGGGGGC	3728
571	CCCCCUG G GGCUGGG	1532	CCGAGCC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CAGGGGG	3729
572	CCCCCUG G GCUCCGG	1533	GCGAGC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG CCAGGGG	3730
578	GGGCGUG G GCUCCCG	1534	GGGGAGC GGAGAAAUCC CU UCAAGGACAUCGUCCGGG GCAGCCCC	3731

Table 23

587	GCUGCCCC G GGAGACCG	1535	CGGUCUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGGGACG	3732
588	CUGCCCCG G GAGACCGA	1536	UCGGUCUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGGGGACG	3733
589	UGCCCCCG G AGACCGAC	1537	GUCGGUCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCGGGGCA	3734
591	CCCCGGGA G ACCGACGA	1538	UCGUCGGU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCCGGGG	3735
601	CCGACGAA G AGCCGAG	1539	CUCGGGCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUCGUCCG	3736
609	GAGCCCGA G GAGCCCGG	1540	CCGGGCUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCGGGCUC	3737
610	AGCCCGAG G AGCCGGC	1541	GCCGGCUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUCGGGCU	3738
616	AGGAGCCC G GCCGAGG	1542	CCUCGGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGGCUCCU	3739
620	GCCGGGCC G GAGGGCA	1543	UGCCCCUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGCCGGGC	3740
621	CCGGGCCG G AGGGCAG	1544	CUGCCCUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGGCCGGG	3741
623	CGGCCGGA G GGGCAGCU	1545	AGCUGCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCGGCCG	3742
624	GGCCGGAG G GGCAGCUU	1546	AAGCUGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUCCGGCC	3743
625	GCCGGAGG G GCAGCUTU	1547	AAAGCUGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCUCGGCC	3744
636	AGCUTUGU G GAGAUGGU	1548	ACCAUCUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACAAGGCU	3745
637	GCUUUGUG G AGAUGGUG	1549	CACCAUCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CACAAAGC	3746
639	UUUGUGGA G AUGGUGGA	1550	UCCACCAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCACAAA	3747
642	GUGGAGAU G GUGACAAA	1551	UUGUCCAC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUCUCCAC	3748
645	GAGAUGGU G GACAACCU	1552	AGGUUGUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACCAUCUC	3749
646	AGAUGGUG G ACAACCTUG	1553	CAGGUUGU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CACCAUCU	3750
656	CAACCUGA G GGGCAAGU	1554	ACUUGCCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCAGGUUG	3751
657	AACCTUGAG G GGCAAGUC	1555	GACUUGCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUCAGGUU	3752
658	ACCUGAGG G GCAAGUCG	1556	CGACUUGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCUCAGGU	3753
666	GGCAAGUC G GGGCAGGG	1557	CCCUGCCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GACUUGCC	3754
667	GCAAGUCG G GGCAGGGC	1558	GCCCUGCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGACUUGC	3755
668	CAAGUCGG G GCAGGGCU	1559	AGCCUUGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCGACUUG	3756
672	UUGGGGCA G GGCUACUA	1560	UAGUAGCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCCCCGA	3757
673	CGGGGCAG G GAGUAC	1561	GUAGUAGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGCCCCG	3758
684	UACUACGU G GAGUAGC	1562	GUCAUCUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACGUAGUA	3759
685	ACTUACUG G AGAUGACC	1563	GGUCAUCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CACGUAGU	3760
687	UACGUGGA G AUGACCGU	1564	ACGUCAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCACGUA	3761
696	AUGACCGU G GGCAGCCC	1565	GGGCUGCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACGGUCAU	3762
697	UGACCGUG G GCAGCCCC	1566	GGGGCUGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CACGUAUA	3763
711	CCCCCGCA G ACGCUCAA	1567	UUGAGCGU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCGGGGG	3764
726	AACAUCU G GUGGAUAC	1568	GUAUCCAC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGGAUGUU	3765

Table 23

729	AUCCUGGU G GAUACAGG	1569	CCUGUAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG ACCAGGAU	3766
730	UCCUGGUG G AUACAGGC	1570	GCCUGUAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CACCAGGA	3767
736	UGGAUACA G GCAGCAGU	1571	ACUGCUGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGUAUCCA	3768
756	UUUGCAGU G GUGUCUGC	1572	GCAGCACC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG ACUGCAAA	3769
757	UUGCAGUG G GUGCUGCC	1573	GGAGCAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CACUGCAA	3770
795	UACUACCA G AGGCAGCU	1574	AGCUGCCU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGUUAGUA	3771
797	CUACCCAGA G GCAGCUGU	1575	ACAGCUGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UCUGGUAG	3772
818	CACAUACC G GCACCUCC	1576	GGAGUCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGUUUGUG	3773
819	ACAUACCG G GACCUCGG	1577	CGGAGUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CGGUUUGU	3774
820	CAUACCGG G ACCUCGGG	1578	CCGAGGUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCGUUAUG	3775
827	GGACCUCC G GAAGGGUG	1579	CACCCUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGAGGUCC	3776
828	GACCUCCG G AAGGGUGU	1580	ACACCCUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CGGAGGUC	3777
831	CUCCGGAA G GUGUGUAU	1581	UACACAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUCGCGAG	3778
832	UCCGGAAG G GUGUGUAU	1582	AUACACAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CUUCCGGA	3779
855	UACACCCA G GCAAGUG	1583	CACUUGCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGGUGUA	3780
856	ACACCCAG G GCAAGUGG	1584	CCACUUGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CUGGGUGU	3781
863	GGCAAGU G GGAGGGG	1585	CCCCUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG ACUUGCCC	3782
864	GGCAAGUG G GAAGGGGA	1586	UCCCCUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CACUUGCC	3783
865	GCAAGUGG G AAGGGGAG	1587	CUCCCCUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCACUUGC	3784
868	AGUGGGAA G GGAGCUG	1588	CAGCUCCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUCCACU	3785
869	GUGGGAG G GGAGCUGG	1589	CCAGCUCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CUUCCAC	3786
870	UGGGAAGG G GAGCUGG	1590	CCCAGCUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCUCCCA	3787
871	GGGAAGG G AGCUGGC	1591	GCCCAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCUUCGCC	3788
876	GGGAGCU G GCACCGA	1592	UCGGUCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGCUCCCC	3789
877	GGGAGCUG G GCACCGAC	1593	GUCGUGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAGCUCC	3790
888	ACGACCU G GUAAGAU	1594	AUGCUUAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGGUUGU	3791
904	UCCCCCAU G GCCCAAC	1595	GUUGGGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUGGGGA	3792
952	CUGAAUCA G ACAAGUUC	1596	GAACUUGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGAUUCAG	3793
970	UCAUCAAC G GCUCCAAC	1597	GUUGGAGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUUGAUGA	3794
980	CUCCAACU G GGAAGGCA	1598	UGCCUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGUUGGAG	3795
981	UCCAACUG G GAAGGCAU	1599	AUGCCUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAGUUGGA	3796
982	CCAACUGG G AAGGCAUC	1600	GAUGCCUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCAGUUGG	3797
985	ACUGGGAA G GCAUCCUG	1601	CAGGAUGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUCGCCAGU	3798
993	GGCAUCCU G GGGCUGGC	1602	GCCAGCCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGGAUGCC	3799

Table 23

994	GCAUCCUG G GCGUGGCC	1603	GGCCAGCC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CAGGAUGC	3800
995	CAUCCUGG G GCUGGCCU	1604	AGGCCAGC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CCAGGAUG	3801
999	CUGGGGU G GCUAUGC	1605	GCAVAGC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AGCCCCAG	3802
1011	UAUGCUGA G AUUGCCAG	1606	CUGGCAU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UCAGCAUA	3803
1019	GAUUGCCA G GCUAGACG	1607	CGUCAGC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UGGCAAUC	3804
1035	GACUCCU G GAGCCUUU	1608	AAAGGCU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AGGAGUC	3805
1036	ACUCCUG G AGCCUUUC	1609	GAAAGCU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CAGGGAGU	3806
1056	GACUCUCU G GUAAAGCA	1610	UGCUUAC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AGAGAGUC	3807
1065	GUAAAGCA G ACCCAGCU	1611	ACUGGGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UGCUUUAC	3808
1102	AGCUUGU G GUGCUGGC	1612	GCCAGCAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG ACAAGCU	3809
1108	GUGGUGU G GCUUCCCC	1613	GGGAAAGC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AGCACCAC	3810
1137	GAAGUGU G GCUUCUGU	1614	ACAGAGC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AGCACUUC	3811
1147	CCUCUGU G GAGGGAGC	1615	GUCCCUU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG GACAGAG	3812
1148	CUCUGUG G AGGGAGCA	1616	UGUCCCU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CGACAGAG	3813
1150	CUGUCGA G GGAGCAUG	1617	CAUGCUC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UCCGACAG	3814
1151	UGUCGGAG G GAGCAUGA	1618	UAUGCUC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CUCCGACA	3815
1152	GUCGGAGG G AGCAUGAU	1619	AUCAUGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CCUCCGAC	3816
1165	UGAUCAUU G GAGUAUC	1620	GAUACCU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AAUGAUA	3817
1166	GAUCAUUG G AGGUAUCG	1621	CGAUACCU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CAAUGAUC	3818
1168	UCAUUGGA G GUUUCGAC	1622	GUCAUAC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UCCAAUGA	3819
1192	UGUACACA G GCAGUCUC	1623	GAGACUGC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UGUGUACA	3820
1202	CAGUCUCU G GUUACAC	1624	GUGUAUAC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AGAGACUG	3821
1217	ACCCAUC G GCGGGAGU	1625	ACUCCGC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG GGAUGGU	3822
1220	CAUCCGC G GGAGUGU	1626	ACCACUC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG GCGGAUG	3823
1221	AUCCGGG G GAGUGUA	1627	UACCACUC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CGCGGAU	3824
1222	UCCGGCG G AGUGGUU	1628	AUACCACU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CCGCCGGA	3825
1226	GCGGGAGU G GUUUAUG	1629	CAUAAUAC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG ACUCCCGC	3826
1236	UAUUAUGA G GUGAUCAU	1630	AUGAUCAC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UCAUUAUA	3827
1250	CAUUGUC G GUGGAGA	1631	UCUCCACC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG GCACAAUG	3828
1251	AUUGUGG G GUGGAGAU	1632	AUCUCCAC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CGCACAAU	3829
1254	GUGCGGGU G GAGAUCAA	1633	UUGAUCUC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG ACCGCAC	3830
1255	UGCGGGUG G AGAUCAAU	1634	AUUGAUCU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CACCGCA	3831
1257	CGGGUGGA G AUCAAUGG	1635	CCAUUGAU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UCCACCGG	3832
1264	AGAUCAAU G GACAGGAU	1636	AUCCUGUC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AUUGAUCU	3833

Table 23

1265	GAUCAAUG G ACAGGAUC	1637	GAUCCUGU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CAUUGAUC	3834
1269	AUUGGACA G GAUCUGAA	1638	UUCAGAUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGUCCAUC	3835
1270	AUGGACAG G AUCUGAAA	1639	UUUCAGAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CUGUCCAUC	3836
1281	CUGAAAAU G GACUGCAA	1640	UUGCAGUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUUUUCAG	3837
1282	UGAAAAUG G ACUGCAAG	1641	UUUGCAGU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CAUUUUCA	3838
1290	GACUGCAA G GAGUACAA	1642	UUUGACUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UUGCAGUC	3839
1291	ACUGCAAG G AGUACAAC	1643	UUUGUACU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CUUGCAGU	3840
1308	UAUGACAA G AGCAUUGU	1644	ACAAUGCU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UUGUCAUA	3841
1317	AGCAUUGU G GACAGUGG	1645	CCACUGUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACAAUUCU	3842
1318	GCAUUGU G ACAGUGGC	1646	GCCACUGU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CACAUAUC	3843
1324	UGGACAGU G GCACCACC	1647	GGUGGUGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACUGUCCA	3844
1350	UUGCCCAA G AAAGUGUU	1648	AACACUUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UUGGGCAA	3845
1383	UCCAUCAA G GCAGCCUC	1649	GAGGUGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UUGAUGGA	3846
1398	UCCUCCAC G GAGAAGUU	1650	AACUUUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG GUGGAGGA	3847
1399	CCUCCACG G AGAAGUUC	1651	GAACUUUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CGUGGAGG	3848
1401	UCCACGGA G AAGUUCU	1652	GGGAACUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UCCGUGGA	3849
1414	UCCUGAU G GUUUCUGG	1653	CCAGAAAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUCAGGGA	3850
1421	UGGUUUUC G GCUAGGAG	1654	CUCUAGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGAAACCA	3851
1426	UCUGGCUA G GAGAGCAG	1655	CUGUCUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UAGCCAGA	3852
1427	CUGGCUAG G AGAGCAGC	1656	GCUCUCU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CUGAGCCAG	3853
1429	GGCUAGGA G AGCAGCUG	1657	CAGUCUCU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UCCUAGCC	3854
1437	GAGCAGCU G GUGUGCUG	1658	CAGCACAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGCUCUCU	3855
1445	GGUGUGCU G GCAAGCAG	1659	CUGCUUGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGCACACC	3856
1453	GGCAGCA G GCACCACC	1660	GGUGGUGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGCUUGCC	3857
1466	CACCCUUU G GAACAUTU	1661	AAAUUUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAGGGUG	3858
1467	ACCCUUG G AACAUUU	1662	AAAUGUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CAAGGGGU	3859
1500	UACCUAAU G GGUGAGGU	1663	ACCUCACC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUUAGGUA	3860
1501	ACCUAAUG G GUGAGGUU	1664	AACCUCAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CAUUAGGU	3861
1506	AUGGGUGA G GUUACCAA	1665	UUUGUAAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UCACCCAU	3862
1556	AUACCUGC G GCCAGUGG	1666	CCACUGGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG GCAGGUUU	3863
1563	CGGCCAGU G GAAAGUUG	1667	ACAUCUUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACUGGCCG	3864
1564	GGCCAGUG G AAGAUGUG	1668	CACAUCUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CACUGGCC	3865
1567	CAGUGGAA G AUGUGGCC	1669	GGCACAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UUCACACUG	3866
1572	GAAGAUGU G GCCACGUC	1670	GACGUGGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACAUCUUC	3867

Table 23

1585	CGUCCAA G ACGACUGU	1671	ACAGUCGU GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG UUGGGACG	3868
1623	UCAUCCAC G GGCACUGU	1672	ACAGUGCC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG GUGGAUGA	3869
1624	CAUCCACG G GCACUGUU	1673	AACAGUGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG CGUGGAUG	3870
1635	ACUGUUU G GGAGCUGU	1674	ACAGCUC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG AUAACAGU	3871
1636	CUGUUU G GAGCUGUU	1675	AACAGCUC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG CAUAACAG	3872
1637	UGUUUUG G AGCUGUUA	1676	UAACAGCU GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG CCAUAACA	3873
1650	GUUAUCAU G GAGGGCUU	1677	AAGCCUC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG AUGAUAA	3874
1651	UUUAUCAU G AGGGCUUC	1678	GAAGCCCU GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG CAUGAUA	3875
1653	AUCAUGGA G GGUUCUA	1679	UAGAAGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG UCCAUGAU	3876
1654	UCAUGGAG G GCUUCUAC	1680	GUAGAAGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG CUCCAUGA	3877
1676	CUUUGAUC G GGCCCGAA	1681	UUCGGGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG GAUCAAG	3878
1677	UUUGAUC G GCUUUGCU	1682	UUUCGGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG GAUCAAA	3879
1693	AACGAAU G GCUUUGCU	1683	AGCAAAGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG AAUUCGUU	3880
1733	UGAGUUA G GACGGCAG	1684	CUGCCGUC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG UGAAACUA	3881
1734	GAGUUCAG G ACGGCAGC	1685	GUCCGCU GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG CUGAACUC	3882
1737	UUCAGGAC G GCAGCGGU	1686	ACCGCUC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG GUCCUGAA	3883
1743	ACGGCAGC G GUGGAAGG	1687	CCUCCAC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG GCUGCCGU	3884
1746	GCAGCGGU G GAAGGCC	1688	GGGCCUUC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG ACCGUCG	3885
1747	CAGCGGUG G AAGGCCU	1689	AGGGCCU GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG CACCGCUG	3886
1750	CGGUGGAA G GCCUUUU	1690	AAAAGGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG UUCCACCG	3887
1767	GUACCUU G GACAUGGA	1691	UCCAUGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG AAGGUGAC	3888
1768	UACCUUG G ACAUGGAA	1692	UCCAUGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG CAAGGUGA	3889
1773	UUGGACAU G GAAGACUG	1693	CAGUCUUC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG AUGUCCAA	3890
1774	UGGACAU G AAGACUGU	1694	ACAGUCU GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG CAUGUCCA	3891
1777	ACAUGGAA G ACUGGGC	1695	GCCACAGU GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG UUCCAUGU	3892
1783	AAGACUGU G GCUACAC	1696	GUUGAGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG ACAGUCUU	3893
1800	AUCCACA G ACAGAUA	1697	UCAUCUGU GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG UGUGGAAU	3894
1804	CACAGACA G AUGAGUCA	1698	UGACUCAU GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG UGUCUGUG	3895
1839	UAUGUCAU G GCUGCCAU	1699	AUGGCAGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG AUGACAUA	3896
1881	UGCCUCAU G GUGUGUA	1700	UGACACAC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG AUGAGGCA	3897
1892	GUGUCAGU G GCGUGCC	1701	GGCAGCGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG ACUGACAC	3898
1960	UGAAGUGA G GAGGCCA	1702	UGGGCCUC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG UCACUUCA	3899
1961	GAAGUGAG G AGGCCAU	1703	AUGGGCCU GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG CUCACUUC	3900
1963	AGUGAGGA G GCCCAUGG	1704	CCAUGGGC GGAGAAAACUCC CU UCAAGGACAUUCGUCCGGG UCCUCACU	3901

Table 23

1970	AGGCCCAU G GGCAGAAG	1705	CUUCUGCC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUGGSCCU	3902
1971	GGCCCAUG G GCAGAAGA	1706	UCUCUGC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG CAUGGGCC	3903
1975	CAUGGGCA G AAGAUAGA	1707	UCUAUCUU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UGCCCAUG	3904
1978	GGCAGAA G AUGAGAU	1708	AUCUCUAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UUCUGCCC	3905
1982	AGAAGUA G AGAUUCCC	1709	GGGAUCU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UAUCUUCU	3906
1984	AAGAUAGA G AUUCCCCU	1710	AGGGAAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UCUAUCUU	3907
1993	AUCCCCU G GACCACAC	1711	GUUGGUC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGGGAAU	3908
1994	UCCCCUG G ACCACACC	1712	GGUGGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG CAGGGAA	3909
2008	ACCUCGU G GUUCACUU	1713	AAGUAA GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG ACGGAGU	3910
2018	UUCACUUU G GUCACAAG	1714	CUUGAC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AAAGUGAA	3911
2029	CACAAGUA G GAGACACA	1715	UGUGUC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UACUUGU	3912
2030	ACAAGUAG G AGACACAG	1716	CUGUCU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG CUACUUGU	3913
2032	AAGUAGGA G ACACAGAU	1717	AUCUGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UCCUACUU	3914
2038	GAGACACA G AUGGCACC	1718	GGUGCAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UGUGUCU	3915
2041	ACACAGAU G GCACUUGU	1719	ACAGGUC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUCUGUGU	3916
2050	GCACCUU G GGCAGAGC	1720	GCUCGUC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG ACAGGUC	3917
2055	UGUGGCA G AGCACCU	1721	GAGGUCU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UGGCACA	3918
2065	GCACCUCA G GACCCUCC	1722	GGAGGUC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UGAGGUC	3919
2066	CACCUAG G ACCUCCC	1723	GGAGGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG CUGAGUG	3920
2101	GCUUGAU G GAGAAGGA	1724	UCCUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUCAGGC	3921
2102	CCUUGAU G AGAAGGAA	1725	UCCUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG CAUCAAG	3922
2104	UUGAUGA G AAGGAAA	1726	UUUCCUU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UCCAUA	3923
2107	AUGGAGAA G GAAAAGGC	1727	GCCUUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UUCUCA	3924
2108	UGGAGAAG G AAAAGGCU	1728	AGCCUUU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG CUUCUCA	3925
2113	AAGGAAA G GCUGGCAA	1729	UUGCCAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UUUUCCUU	3926
2117	AAAAGGCU G GUGGUUC	1730	CACUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGCCUUUU	3927
2122	GCUGGCA G GUGGUUC	1731	GAACCA GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UUGCCAG	3928
2125	GGCAAGU G GUUCCAG	1732	CUGAAC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG ACCUUGC	3929
2126	GCAAGGU G GUUCCAG	1733	CCUGAAC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG CACCUUG	3930
2133	GGUUGCA G GGACUUA	1734	UACAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UGGAACCC	3931
2134	GGUUGCA G GACUUA	1735	GUACAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG CUGGAACC	3932
2135	GUUCCAG G ACUGUAC	1736	GGUACAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG CCUGAAC	3933
2148	UACCUAG G GAAACAGA	1737	UCUGUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UACAGUA	3934
2149	ACCUGUAG G AAACAGA	1738	UUCUGUU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG CUACAGU	3935

Table 23

2155	AGGAACA G AAAAGAGA	1739	UCUCUUU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UGUUUCU	3936
2160	ACAGAAA G AGAAGAA	1740	UUUCUUU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UUUUUCU	3937
2162	AGAAAGA G AAGAAAG	1741	UCUUUUU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UCUUUUU	3938
2165	AAAGAGAA G AAAGAAGC	1742	GCUUUUU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UUCUCUU	3939
2169	AGAGAGAA G AAGCACUC	1743	GAGUGUU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UUUUUUU	3940
2182	ACUCUGU G GCGGAAU	1744	AUCCCCG GGAGGAAACUCC CU UCAAGGACAUCGUCGGG AGCAGAGU	3941
2185	CUGCUGG G GGAUACU	1745	AGAUUCC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG GCCAGCAG	3942
2186	UGCUGGG G GAUACUC	1746	GAGAUUC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG CGCCAGCA	3943
2187	GCUGGGG G AAUACUCU	1747	AGAGUUU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG CCGCCAGC	3944
2197	AUACUCU G GUCACCUC	1748	GAGGUGA GGAGGAAACUCC CU UCAAGGACAUCGUCGGG AAGAGUUA	3945
2217	UUUAAGUC G GGAUUUC	1749	GAAUUCC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG GACUUAAA	3946
2218	UUAAGUCG G GAAUUCU	1750	AGAAUUC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG CGACUUAA	3947
2219	UAAGUCGG G AAUUCUG	1751	CAGAUUU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG CCGACUUA	3948
2311	UAGUUUA G AAGUACUG	1752	CAGUACU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UGAAACUA	3949
2319	GAAGUACU G GCAUCACA	1753	UGUGAUC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG AGUACUUC	3950
2332	CACAGCA G GUUACCUU	1754	AAGGUAC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UGCGUGUG	3951
2341	GUUACCU G GCGUGUGU	1755	ACACAGC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG AAGGUAA	3952
2356	GUCCUCU G GUACCCUG	1756	CAGGUAC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG ACAGGGAC	3953
2364	GGUACCCU G GCAGAGAA	1757	UUUCUCU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG AGGGUACC	3954
2368	CCUUGGA G AGAAGAGA	1758	UCUUUCU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UGCCAGGG	3955
2370	CUGGCAG G AAGAGACC	1759	GGUCUCU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UGUCGCCAG	3956
2373	GCAGAGAA G AGACCAAG	1760	CUUGGUC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UUCUCUGC	3957
2375	AGAGAGAA G ACCAAGCU	1761	AGCUUGU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UCUUCUCU	3958
2396	UCCUCUGU G GCCAAAGU	1762	ACUUUGC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG AGCAGGGA	3959
2410	AGUCAGUA G GAGAGGAU	1763	AUCCUCU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UACUGACU	3960
2411	GUCAGUAG G AGAGGAUG	1764	CAUCCUC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG CUACUGAC	3961
2413	CAGUAGGA G AGGAUGCA	1765	UGCAUCCU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UCCUACUG	3962
2415	GUAGGAGA G GAUGCACA	1766	UGUGAUC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UCUCCUAC	3963
2416	UAGGAGAG G AUGCACAG	1767	CUGUGCAU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG CUUCCUUA	3964
2441	UUGCUUUA G AGACAGGG	1768	CCUUGUCU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UAAAGCAA	3965
2443	GCUUUAGA G ACAGGGAC	1769	GUCCUGU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UCUAAAGC	3966
2447	UAGAGACA G GGACUGUA	1770	UACAGUCC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG UGUUCUUA	3967
2448	AGAGACAG G GACUGUAU	1771	AUACAGUC GGAGGAAACUCC CU UCAAGGACAUCGUCGGG CUGUCUCU	3968
2449	GAGACAG G ACUGUAUA	1772	UAUACAGU GGAGGAAACUCC CU UCAAGGACAUCGUCGGG CCUGUCUC	3969

Table 23

2473	CUAACAUU G GUGCAAAG	1773	CUUUGCAC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AAUGUUAG	3970
2481	GGUGCAAA G AUGCCUC	1774	GAGGCAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UUUGCACC	3971
2511	AAAAACUA G AAAAAAA	1775	UUUUUUU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG UAGUUUUU	3972

Input Sequence = AF190725. Cut Site = G/.
Stem Length = 8. Core Sequence = GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG
AF190725 (Homo sapiens beta-site APP cleaving enzyme (BACE) mRNA; 2526 bp)

Table 24

Table 24: Human Phospholamban (PLN) Hammerhead Ribozyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
16	AGAAAACU C CCCAGCUA	1	UAGCUGGG CUGAUGAG X CGAA AGUUUUCU	1137
24	CCCCAGCU A AACACCCG	2	CGGGUGUU CUGAUGAG X CGAA AGCUGGGG	1138
34	ACACCCGU A AGACUUC A	3	UGAAGUCU CUGAUGAG X CGAA ACGGGUGU	1139
40	GUAAGACU U CAUACAAC	4	GUUGUAUG CUGAUGAG X CGAA AGUCUUC	1140
41	UAAGACUU C AUACAACA	5	UGUUGUAU CUGAUGAG X CGAA AAGUCUUA	1141
44	GACUUCAU A CAACACAA	6	UUGUGUUG CUGAUGAG X CGAA AUGAAGUC	1142
54	AACACAAU A CUCUAUAC	7	GUUAUGAG CUGAUGAG X CGAA AUUGUGUU	1143
57	ACAAUACU C UAUACUGU	8	ACAGUAUA CUGAUGAG X CGAA AGUAUUGU	1144
59	AAUACUCU A UACUGUGA	9	UCACAGUA CUGAUGAG X CGAA AGAGUAUU	1145
61	UACUCUAU A CUGUGAUG	10	CAUCACAG CUGAUGAG X CGAA AUAGAGUA	1146
72	GUGAUGAU C ACAGCUGC	11	GCAGCUGU CUGAUGAG X CGAA AUCAUCAC	1147
88	CCAAGGCU A CCUAAAAG	12	CUUUUAGG CUGAUGAG X CGAA AGCCUUGG	1148
92	GGCUACCU A AAAGAAGA	13	UCUUCUUU CUGAUGAG X CGAA AGGUAGCC	1149
105	AAGACAGU U AUCUCAUA	14	UAUGAGAU CUGAUGAG X CGAA ACUGUCUU	1150
106	AGACAGUU A UCUCUAUU	15	AUAUGAGA CUGAUGAG X CGAA AACUGUCU	1151
108	ACAGUUAU C UCAUAUUU	16	AAAUAUGA CUGAUGAG X CGAA AUAACUGU	1152
110	AGUUAUCU C AUAUUUGG	17	CCAAUAUU CUGAUGAG X CGAA AGAUAACU	1153
113	UAUCUCAU A UUUGGCUG	18	CAGCCAAA CUGAUGAG X CGAA AUGAGUAU	1154
115	UCUCAUAU U UGGCUGCC	19	GGCAGCCA CUGAUGAG X CGAA AUAUGAGA	1155
116	CUCAUAUU U GGCUGCCA	20	UGGCAGCC CUGAUGAG X CGAA AAUAUGAG	1156
128	UGCCAGCU U UUAUCUU	21	AAGAUAAA CUGAUGAG X CGAA AGCUGGCA	1157
129	GCCAGCUU U UUAUCUU	22	AAAGAUAA CUGAUGAG X CGAA AAGCUGGC	1158
130	CCAGCUUU U UAUCUUUC	23	GAAAGAU A CUGAUGAG X CGAA AAAGCUGG	1159
131	CAGCUUUU U AUCUUUCU	24	AGAAAGAU CUGAUGAG X CGAA AAAAGCUG	1160
132	AGCUUUUU A UCUUUCUC	25	GAGAAAGA CUGAUGAG X CGAA AAAAAGCU	1161
134	CUUUUUAU C UUUCUCUC	26	GAGAGAAA CUGAUGAG X CGAA AUAAAAAG	1162
136	UUUUUAUCU U UCUCUCGA	27	UCGAGAGA CUGAUGAG X CGAA AGAUAAAA	1163
137	UUUAUCUU U CUCUCGAC	28	GUCGAGAG CUGAUGAG X CGAA AAGAUAAA	1164
138	UUAUCUUU C UCUCGACC	29	GGUCGAGA CUGAUGAG X CGAA AAAGAUAA	1165
140	AUCUUUCU C UCGACCAC	30	GUGGUCGA CUGAUGAG X CGAA AGAAAGAU	1166
142	CUUUCUCU C GACCACUU	31	AAGUGGUC CUGAUGAG X CGAA AGAGAAAG	1167
150	CGACCACU U AAAACUUC	32	GAAGUUUU CUGAUGAG X CGAA AGUGGUCG	1168
151	GACCACUU A AAACUUCA	33	UGAAGUUU CUGAUGAG X CGAA AAGUGGUC	1169
157	UUAAAACU U CAGACUUC	34	GAAGUCUG CUGAUGAG X CGAA AGUUUUAA	1170
158	UAAAACUU C AGACUUC	35	GGAAGUCU CUGAUGAG X CGAA AAGUUUUA	1171
164	UUCAGACU U CCUGUCCU	36	AGGACAGG CUGAUGAG X CGAA AGUCUGAA	1172
165	UCAGACUU C CUGUCCUG	37	CAGGACAG CUGAUGAG X CGAA AAGUCUGA	1173
170	CUUCCUGU C CUGCUGGU	38	ACCAGCAG CUGAUGAG X CGAA ACAGGAAG	1174
179	CUGCUGGU A UCAUGGAG	39	CUCCAUGA CUGAUGAG X CGAA ACCAGCAG	1175
181	GCUGGUAU C AUGGAGAA	40	UUCUCCAU CUGAUGAG X CGAA AUACCAGC	1176
193	GAGAAAGU C CAAUACCU	41	AGGUUAUG CUGAUGAG X CGAA ACUUUCUC	1177
198	AGUCCAAU A CCUCACUC	42	GAGUGAGG CUGAUGAG X CGAA AUUGGACU	1178

Table 24

202	CAAUACCU C ACUCGCUC	43	GAGCGAGU CUGAUGAG X CGAA AGGUAUUG	1179
206	ACCUCACU C GCUCAGCU	44	AGCUGAGC CUGAUGAG X CGAA AGUGAGGU	1180
210	CACUCGCU C AGCUAUA	45	UUAUAGCU CUGAUGAG X CGAA AGCGAGUG	1181
215	GCUCAGCU A UAAGAAGA	46	UCUUCUUA CUGAUGAG X CGAA AGCUGAGC	1182
217	UCAGCUAU A AGAAGAGC	47	GCUCUUCU CUGAUGAG X CGAA AUAGCUGA	1183
228	AAGAGCCU C AACCAUUG	48	CAAUGGUU CUGAUGAG X CGAA AGGCUCUU	1184
235	UCAACCAU U GAAAUGCC	49	GGCAUUTC CUGAUGAG X CGAA AUGGUUGA	1185
245	AAAUGCCU C AACCAAGCA	50	UGCUGUUU CUGAUGAG X CGAA AGGCAUUU	1186
257	AAGCACGU C AAAAGCUA	51	UAGCUUUU CUGAUGAG X CGAA ACGUGCUU	1187
265	CAAAGCCU A CAGAAUCU	52	AGAUCUG CUGAUGAG X CGAA AGCUUUUG	1188
272	UACAGAAU C UAUUUUUC	53	GAUAAUA CUGAUGAG X CGAA AUUCUGUA	1189
274	CAGAAUCU A UUUUAUAA	54	UUGAUAAA CUGAUGAG X CGAA AGAUUCUG	1190
276	GAAUCUAU U UAUCAAUU	55	AAUUGAUA CUGAUGAG X CGAA AUAGAUUC	1191
277	AAUCUAUU U AUCAAUUU	56	AAAUUGAU CUGAUGAG X CGAA AAUAGAUU	1192
278	AUCUAUUU A UCAAUUUC	57	GAAAUUGA CUGAUGAG X CGAA AAUAGAU	1193
280	CUAUUUU C AAUUCUG	58	CAGAAAUU CUGAUGAG X CGAA AUAAUAG	1194
284	UUAUCAAU U UCUGUCUC	59	GAGACAGA CUGAUGAG X CGAA AUUGAUAA	1195
285	UAUCAAUU U CUGUCUCA	60	UGAGACAG CUGAUGAG X CGAA AAUUGAUA	1196
286	AUCAAUUU C UGUCUCAU	61	AUGAGACA CUGAUGAG X CGAA AAUUGAU	1197
290	AUUUCUGU C UCAUCUUA	62	UAAGAUGA CUGAUGAG X CGAA ACAGAAAU	1198
292	UUCUGUCU C AUCUUAU	63	AUUAAGAU CUGAUGAG X CGAA AGACAGAA	1199
295	UGUCUCAU C UUAUAUG	64	CAUAUUAA CUGAUGAG X CGAA AUGAGACA	1200
297	UCUCAUCU U AAUAUGUC	65	GACAUUU CUGAUGAG X CGAA AGAUGAGA	1201
298	CUCAUCUU A AUAUGUCU	66	AGACAUUU CUGAUGAG X CGAA AAGAUGAG	1202
301	AUCUUAU A UGUCUCUU	67	AAGAGACA CUGAUGAG X CGAA AUUAAGAU	1203
305	UAUAUGU C UCUGCUG	68	CAGCAAGA CUGAUGAG X CGAA ACAUAUUA	1204
307	AUAUGUCU C UUGCUGAU	69	AUCAGCAA CUGAUGAG X CGAA AGACAUUU	1205
309	AUGUCUCU U GCUGAUCU	70	AGAUCAGC CUGAUGAG X CGAA AGAGACAU	1206
316	UUGCUGAU C UGUUAUUA	71	AUGAUACA CUGAUGAG X CGAA AUCAGCAA	1207
320	UGAUCUGU A UCAUCGUG	72	CACGAUGA CUGAUGAG X CGAA ACAGAUCA	1208
322	AUCUGUAU C AUCGUGAU	73	AUCACGAU CUGAUGAG X CGAA AUACAGAU	1209
325	UGUAUCAU C GUGAUGCU	74	AGCAUCAC CUGAUGAG X CGAA AUGAUACA	1210
334	GUGAUGCU U CUCUGAAG	75	CUUCAGAG CUGAUGAG X CGAA AGCAUCAC	1211
335	UGAUGCUU C UCUGAAGU	76	ACUUCAGA CUGAUGAG X CGAA AAGCAUCA	1212
337	AUGCUUCU C UGAAGUUC	77	GAACUUCA CUGAUGAG X CGAA AGAAGCAU	1213
344	UCUGAAGU U CUGCUACA	78	UGUAGCAG CUGAUGAG X CGAA ACUUCAGA	1214
345	CUGAAGUU C UGCUACAA	79	UUGUAGCA CUGAUGAG X CGAA AACUUCAG	1215
350	GUUCUGCU A CAACCUCU	80	AGAGGUUG CUGAUGAG X CGAA AGCAGAAC	1216
357	UACAACCU C UAGAUCUG	81	CAGAUCUA CUGAUGAG X CGAA AGGUUGUA	1217
359	CAACCUCU A GAUCUGCA	82	UGCAGAU CUGAUGAG X CGAA AGAGGUUG	1218
363	CUCUAGAU C UGCAGCUU	83	AAGCUGCA CUGAUGAG X CGAA AUCUAGAG	1219
371	CUGCAGCU U GCCACAUC	84	GAUGUGGC CUGAUGAG X CGAA AGCUGCAG	1220
379	UGCCACAU C AGCUUAAA	85	UUUAAGCU CUGAUGAG X CGAA AUGUGGCA	1221
384	CAUCAGCU U AAAAUCUG	86	CAGAUUUU CUGAUGAG X CGAA AGCUGAUG	1222
385	AUCAGCUU A AAAUCUGU	87	ACAGAUUU CUGAUGAG X CGAA AAGCUGAU	1223
390	CUUAAAAU C UGUCAUCC	88	GGAUGACA CUGAUGAG X CGAA AUUUUAG	1224
394	AAAUCUGU C AUCCCAUG	89	CAUGGGAU CUGAUGAG X CGAA ACAGAUUU	1225

Table 24

397	UCUGUCAU C CCAUGCAG	90	CUGCAUGG CUGAUGAG X CGAA AUGACAGA	1226
419	AAAACAAU A UUGUAUAA	91	UUAUACAA CUGAUGAG X CGAA AUUGUUUU	1227
421	AACAAUUAU U GUAUAACA	92	UGUUAUAC CUGAUGAG X CGAA AUUUGUUU	1228
424	AAUAUUGU A UAACAGAC	93	GUCUGUUA CUGAUGAG X CGAA ACAUAUUU	1229
426	UAUUGUAU A ACAGACCA	94	UGGUCUGU CUGAUGAG X CGAA AUACAAUA	1230
437	AGACCACU U CCUGAGUA	95	UACUCAGG CUGAUGAG X CGAA AGUGGUCU	1231
438	GACCACUU C CUGAGUAG	96	CUACUCAG CUGAUGAG X CGAA AAGUGGUC	1232
445	UCCUGAGU A GAAGAGUU	97	AACUCUUC CUGAUGAG X CGAA ACUCAGGA	1233
453	AGAAGAGU U UCUUUGUG	98	CACAAAGA CUGAUGAG X CGAA ACUCUUCU	1234
454	GAAGAGUU U CUUUGUGA	99	UCACAAAG CUGAUGAG X CGAA AACUCUUC	1235
455	AAGAGUUU C UUUGUGAA	100	UUCACAAA CUGAUGAG X CGAA AAACUCUU	1236
457	GAGUUUCU U UGUGAAAA	101	UUUUCACA CUGAUGAG X CGAA AGAAACUC	1237
458	AGUUUCUU U GUGAAAAG	102	CUUUUCAC CUGAUGAG X CGAA AAGAAACU	1238
469	GAAAAGGU C AAGAUUAA	103	UUAUUCUU CUGAUGAG X CGAA ACCUUUUC	1239
475	GUCAAGAU U AAGACUAA	104	UUAGUCUU CUGAUGAG X CGAA AUCUUGAC	1240
476	UCAAGAUU A AGACUAAA	105	UUUAGUCU CUGAUGAG X CGAA AAUCUUGA	1241
482	UUAAGACU A AAACUUUA	106	AUAAGUUU CUGAUGAG X CGAA AGUCUUAA	1242
488	CUAAAACU U AUUGUUAC	107	GUAACAAU CUGAUGAG X CGAA AGUUUUAG	1243
489	UAAAACUU A UUGUUACC	108	GGUAACAA CUGAUGAG X CGAA AAGUUUUA	1244
491	AAACUUUAU U GUUACCAU	109	AUGGUUAC CUGAUGAG X CGAA AUAAGUUU	1245
494	CUUAUUGU U ACCAUUAG	110	CAUAUGGU CUGAUGAG X CGAA ACAUAUAG	1246
495	UUAUUGUU A CCAUAUGU	111	ACAUAUGG CUGAUGAG X CGAA AACAUUAA	1247
500	GUUACCAU A UGUUUUCA	112	UGAAUACA CUGAUGAG X CGAA AUGGUUAC	1248
504	CCAUAUGU A UUCAUCUG	113	CAGAUGAA CUGAUGAG X CGAA ACAUAUGG	1249
506	AUAUGUAU U CAUCUGUU	114	AACAGAUG CUGAUGAG X CGAA AUACAUUU	1250
507	UAUGUAUU C AUCUGUUG	115	CAACAGAU CUGAUGAG X CGAA AAUACAUU	1251
510	GUUUAUUAU C UGUUGGAU	116	AUCCAACA CUGAUGAG X CGAA AUGAAUAC	1252
514	UCAUCUGU U GGAUCUUG	117	CAAGAUCU CUGAUGAG X CGAA ACAGAUGA	1253
519	UGUUGGAU C UUGUAAAC	118	GUUUACAA CUGAUGAG X CGAA AUCCAACA	1254
521	UUGGAUCU U GUAAACAU	119	AUGUUUAC CUGAUGAG X CGAA AGAUCCAA	1255
524	GAUCUUGU A AACAUCAA	120	UUAUUGUU CUGAUGAG X CGAA ACAAGUUC	1256
540	AAAGGGCU U UAUUUUCA	121	UGAAAAUA CUGAUGAG X CGAA AGCCUUUU	1257
541	AAGGGCUU U AUUUUCAA	122	UUGAAAAU CUGAUGAG X CGAA AAGCCUUU	1258
542	AGGGCUUU A UUUUCAA	123	UUUGAAAA CUGAUGAG X CGAA AAAGCCUU	1259
544	GGCUUUUAU U UUCAAAAA	124	UUUUUGAA CUGAUGAG X CGAA AUAAAGCC	1260
545	GCUUUAUU U UCAAAAAU	125	AUUUUUGA CUGAUGAG X CGAA AAUAAAGC	1261
546	CUUUUAUU U CAAAAAUU	126	AAUUUUUG CUGAUGAG X CGAA AAUAAAG	1262
547	UUUAUUUU C AAAAAUUA	127	UAAUUUUU CUGAUGAG X CGAA AAAAAUAA	1263
554	UCAAAAAU U AACUCAA	128	UUGAAGUU CUGAUGAG X CGAA AUUUUUGA	1264
555	CAAAAAUU A ACUCAA	129	UUUGAAGU CUGAUGAG X CGAA AAUUUUUG	1265
559	AAUUAACU U CAAAAUAA	130	UUUUUUUG CUGAUGAG X CGAA AGUUAAUU	1266
560	AUUUAACU C AAAAUUAG	131	CUUUUUUU CUGAUGAG X CGAA AAGUUAAU	1267
566	UUCAAAAU A AGUGUAUA	132	UAUACACU CUGAUGAG X CGAA AUUUUGAA	1268
572	AUAAGUGU A UAAAAUGC	133	GCAUUUUA CUGAUGAG X CGAA ACACUUUU	1269
574	AAGUGUAU A AAAUGCAA	134	UUGCAUUU CUGAUGAG X CGAA AUACACUU	1270
587	GCAACUGU U GAUUUCCU	135	AGGAAUUC CUGAUGAG X CGAA ACAGUUGC	1271
591	CUGUUGAU U UCCUCAAC	136	GUUGAGGA CUGAUGAG X CGAA AUCAACAG	1272

Table 24

592	UGUUGAUU U CCUCAACA	137	UGUUGAGG CUGAUGAG X CGAA AAUCAACA	1273
593	GUUGAUUU C CUCAACAU	138	AUGUUGAG CUGAUGAG X CGAA AAAUCAAC	1274
596	GAUUUCCU C AACAUGGC	139	GCCAUGUU CUGAUGAG X CGAA AGGAAAUC	1275
606	ACAUGGCU C ACAAUUU	140	AAAUUUUU CUGAUGAG X CGAA AGCCAUGU	1276
613	UCACAAAU U UCUAUCCC	141	GGGAUAGA CUGAUGAG X CGAA AUUUGUGA	1277
614	CACAAAUU U CUAUCCCA	142	UGGGAUAG CUGAUGAG X CGAA AAUUGUG	1278
615	ACAAAUUU C UAUCCCAA	143	UUGGGAUA CUGAUGAG X CGAA AAUUGUGU	1279
617	AAAUUUUC A UCCCAAU	144	AUUUGGGA CUGAUGAG X CGAA AGAAAUUU	1280
619	AUUUCUUA C CCAAUCU	145	AGAUAUGG CUGAUGAG X CGAA AUAGAAAU	1281
626	UCCCAAU C UUUUCUGA	146	UCAGAAA CUGAUGAG X CGAA AUUUGGGA	1282
628	CCAAUCU U UUCUGAAG	147	CUUCAGAA CUGAUGAG X CGAA AGAUUUGG	1283
629	CAAAUCU U UCUGAAGA	148	UCUUCAGA CUGAUGAG X CGAA AAGAUUUG	1284
630	AAAUUUU U CUGAAGAU	149	AUCUUCAG CUGAUGAG X CGAA AAAGAUUU	1285
631	AAUCUUU C UGAAGAUG	150	CAUCUUA CUGAUGAG X CGAA AAAAGAUU	1286
646	UGAAGAGU U UAGUUUA	151	UAAAACUA CUGAUGAG X CGAA ACUCUUA	1287
647	GAAGAGUU U AGUUUAA	152	UUAAAACU CUGAUGAG X CGAA AACUCUUC	1288
648	AAGAGUUU A GUUUAAA	153	UUUAAAAC CUGAUGAG X CGAA AAACUCUU	1289
651	AGUUUAGU U UUAAAACU	154	AGUUUUA CUGAUGAG X CGAA ACUAAAACU	1290
652	GUUUAGUU U UAAAACUG	155	CAGUUUA CUGAUGAG X CGAA AACUAAAC	1291
653	UUUAGUUU U AAAACUGC	156	GCAGUUU CUGAUGAG X CGAA AAACUAAA	1292
654	UUAGUUU A AAACUGCA	157	UGCAGUU CUGAUGAG X CGAA AAAACUAA	1293
675	CAACAAGU U CACUUAU	158	AUGAAGUG CUGAUGAG X CGAA ACUUGUUG	1294
676	AACAAGU C ACUUAUA	159	UAUGAAGU CUGAUGAG X CGAA AACUUGUU	1295
680	AGUUCACU U CAUAUAU	160	UAUAUAUG CUGAUGAG X CGAA AGUGAACU	1296
681	GUUCACU C AUAUAUA	161	UUAUAUAU CUGAUGAG X CGAA AAGUGAAC	1297
684	CACUUAU A UAUAAGC	162	GCUUAUA CUGAUGAG X CGAA AUGAAGUG	1298
686	CUUCAUAU A UAAAGCAU	163	AUGCUUA CUGAUGAG X CGAA AUAUGAAG	1299
688	UCAUAUAU A AAGCAUA	164	UAAUGCU CUGAUGAG X CGAA AUAUAUGA	1300
695	UAAAGCAU U AUUUUAC	165	GUAAAAU CUGAUGAG X CGAA AUGCUUA	1301
696	AAAGCAU A UUUUACU	166	AGUAAAA CUGAUGAG X CGAA AAUGCUUU	1302
698	AGCAUAU U UUUACUCU	167	AGAGUAAA CUGAUGAG X CGAA AUAAUGCU	1303
699	GCAUAUAU U UUACUCU	168	AAGAGUAA CUGAUGAG X CGAA AAUAUUGC	1304
700	CAUAUAU U UACUCUU	169	AAAGAGUA CUGAUGAG X CGAA AAAUAUUG	1305
701	AUAUAUUU U ACUCUUU	170	AAAAGAGU CUGAUGAG X CGAA AAAUAUAA	1306
702	UAUAUUU A CUCUUUG	171	CAAAAGAG CUGAUGAG X CGAA AAAUAUA	1307
705	UUUUUACU C UUUUGAGG	172	CCUCAAAA CUGAUGAG X CGAA AGUAAAAA	1308
707	UUUACUCU U UUGAGGUG	173	CACCUCAA CUGAUGAG X CGAA AGAGUAAA	1309
708	UUACUCUU U UGAGGUGA	174	UCACCUC CUGAUGAG X CGAA AAGAGUAA	1310
709	UACUCUUU U GAGGUGAA	175	UUCACCUC CUGAUGAG X CGAA AAAGAGUA	1311
719	AGGUGAAU A UAAUUUAU	176	AUAAAUUA CUGAUGAG X CGAA AUUCACCU	1312
721	GUGAAUAU A AUUAUAU	177	AUAUAAU CUGAUGAG X CGAA AUUAUCAC	1313
724	AAUAUAU U UAUAUAC	178	GUAAUAUA CUGAUGAG X CGAA AUUAUAU	1314
725	AUAUAUU U AUAUAUA	179	UGUAUAU CUGAUGAG X CGAA AAUAUAU	1315
726	UAUAUUU A UAUAACA	180	UUGUAUA CUGAUGAG X CGAA AAUAUAU	1316
728	UAUUUAU A UAUAUAU	181	CAUUGUAA CUGAUGAG X CGAA AUAAUAU	1317
730	AUUUAUAU U ACAAUGUA	182	UACAUUGU CUGAUGAG X CGAA AUUAUAU	1318
731	UUUAUAU A CAAUGUAA	183	UUACAUG CUGAUGAG X CGAA AAUAUAAA	1319

Table 24

738	UACAAUGU A AAAGCUUC	184	GAAGCUUU CUGAUGAG X CGAA ACAUUGUA	1320
745	UAAAAGCU U CUUAAUA	185	UAUUAAAG CUGAUGAG X CGAA AGCUUUUA	1321
746	AAAAGCUU C UUUAAUAC	186	GUUUUAAA CUGAUGAG X CGAA AAGCUUUU	1322
748	AAGCUCU U UAAUACUA	187	UAGUAUUA CUGAUGAG X CGAA AGAAGCUU	1323
749	AGCUUCU U AAUACUAA	188	UUAGUAUU CUGAUGAG X CGAA AAGAAGCU	1324
750	GCUCUUU A AUACUAAAG	189	CUUAGUAU CUGAUGAG X CGAA AAAGAAGC	1325
753	UCUUUAAU A CUAAGUAU	190	AUACUUAG CUGAUGAG X CGAA AUUAAAGA	1326
756	UUAAUACU A AGUAUUUU	191	AAAUAUCU CUGAUGAG X CGAA AGUAUUUA	1327
760	UACUAAGU A UUUUUCAG	192	CUGAAAAA CUGAUGAG X CGAA ACUUAGUA	1328
762	CUAAGUAU U UUUCAGGU	193	ACCUGAAA CUGAUGAG X CGAA AUACUUAG	1329
763	UAAGUAUU U UUCAGGUC	194	GACCUGAA CUGAUGAG X CGAA AAUACUUA	1330
764	AAGUAUUU U UCAGGUCU	195	AGACCUGA CUGAUGAG X CGAA AAAUACUU	1331
765	AGUAUUUU U CAGGUCUU	196	AAGACCUG CUGAUGAG X CGAA AAAAUACU	1332
766	GUUUUUU C AGGUCUUC	197	GAAGACCU CUGAUGAG X CGAA AAAAUAC	1333
771	UUUCAGGU C UUCACCAA	198	UUGGUGAA CUGAUGAG X CGAA ACCUGAAA	1334
773	UCAGGUCU U CACCAAGU	199	ACUUGGUG CUGAUGAG X CGAA AGACCUGA	1335
774	CAGGUCUU C ACCAAGUA	200	UACUUGGU CUGAUGAG X CGAA AAGACCUG	1336
782	CACCAAGU A UCAAAGUA	201	UACUUUGA CUGAUGAG X CGAA ACUUGGUG	1337
784	CCAAGUAU C AAAGUAAU	202	AUUACUUU CUGAUGAG X CGAA AUACUUGG	1338
790	AUCAAGU A AUAACACA	203	UGUGUUUU CUGAUGAG X CGAA ACUUGAU	1339
793	AAAGUAAU A ACACAAU	204	AUUUGUGU CUGAUGAG X CGAA AUUACUUU	1340
809	UGAAGUGU C AUUAUUA	205	UGAAUAAU CUGAUGAG X CGAA ACACUUA	1341
812	AGUGUCAU U AUUCAAU	206	UUUUGAAU CUGAUGAG X CGAA AUGACACU	1342
813	GUGUCAUU A UUCAAAU	207	AUUUUGAA CUGAUGAG X CGAA AAUGACAC	1343
815	GUCAUUU U CAAAUAAG	208	CUAUUUUG CUGAUGAG X CGAA AUAAUGAC	1344
816	UCAUUUU C AAAUAAGU	209	ACUAUUUU CUGAUGAG X CGAA AAUAUGA	1345
822	UUCAAAU A GUCCACUG	210	CAGUGGAC CUGAUGAG X CGAA AUUUUGAA	1346
825	AAAUAAGU C CACUGACU	211	AGUCAGUG CUGAUGAG X CGAA ACUAUUUU	1347
834	CACUGACU C CUCACAUC	212	GAUGUGAG CUGAUGAG X CGAA AGUCAGUG	1348
837	UGACUCCU C ACAUCUGU	213	ACAGAUGU CUGAUGAG X CGAA AGGAGUCA	1349
842	CCUCACAU C UGUUAUCU	214	AGAUAAUA CUGAUGAG X CGAA AUGUGAGG	1350
846	ACAUCUGU U AUUUUAUU	215	AAUAAGAU CUGAUGAG X CGAA ACAGAUGU	1351
847	CAUCUGUU A UCUUUAUA	216	UAUAAGA CUGAUGAG X CGAA AACAGAUG	1352
849	UCUGUUU C UUAUUUA	217	UAUAUAA CUGAUGAG X CGAA AUAACAGA	1353
851	UGUUAUCU U AUUAUAA	218	UUUAUAAU CUGAUGAG X CGAA AGAUAAUA	1354
852	GUUAUCUU A UUAUAAAG	219	CUUAUAA CUGAUGAG X CGAA AAGUAAC	1355
854	UAUCUUU U AUAAAGAA	220	UUCUUUAU CUGAUGAG X CGAA AUAAAGUA	1356
855	AUCUUUU A UAAAGAAC	221	GUUCUUUA CUGAUGAG X CGAA AAUAAGAU	1357
857	CUUAUUU A AAGAACUA	222	UAGUUCUU CUGAUGAG X CGAA AUAAUAAG	1358
865	AAAGAACU A UUGUAAGU	223	ACUACAAA CUGAUGAG X CGAA AGUUCUUU	1359
867	AGAACUAU U UGUAGUA	224	UUACUACA CUGAUGAG X CGAA AUAGUUCU	1360
868	GAACUAUU U GUAGUAAC	225	GUUACUAC CUGAUGAG X CGAA AAUAGUUC	1361
871	CUAUUUGU A GUAACUAU	226	AUAGUUAC CUGAUGAG X CGAA ACAAAUAG	1362
874	UUUGUAGU A ACUAUCAG	227	CUGAUAGU CUGAUGAG X CGAA ACUACAAA	1363
878	UAGUAACU A UCAGAAUC	228	GAUUCUGA CUGAUGAG X CGAA AGUUACUA	1364
880	GUAACUAU C AGAAUCUA	229	UAGAUUCU CUGAUGAG X CGAA AUAGUAC	1365
886	AUCAGAAU C UACAUUCU	230	AGAAUGUA CUGAUGAG X CGAA AUUCUGAU	1366

Table 24

888	CAGAAUCU A CAUUCUAA	231	UUAGAAUG CUGAUGAG X CGAA AGAUUCUG	1367
892	AUCUACAU U CUAAAACA	232	UGUUUUAG CUGAUGAG X CGAA AUGUAGAU	1368
893	UCUACAUU C UAAAACAG	233	CUGUUUUA CUGAUGAG X CGAA AAUGUAGA	1369
895	UACAUCU A AACAGAA	234	UUCUGUUU CUGAUGAG X CGAA AGAAUGUA	1370
906	ACAGAAU U GUUUUUU	235	AAAAAUAC CUGAUGAG X CGAA AUUUCUGU	1371
909	GAAAUUGU A UUUUUUCU	236	AGAAAAA CUGAUGAG X CGAA ACAAUUUC	1372
911	AAUUGUAU U UUUUCUAA	237	AUAGAAAA CUGAUGAG X CGAA AUACAAU	1373
912	AUUGUAU U UUUUCUAG	238	CAUAGAAA CUGAUGAG X CGAA AAUACAAU	1374
913	UUGUAUUU U UUCUAUGC	239	GCAUAGAA CUGAUGAG X CGAA AAAUACAA	1375
914	UGUAUUUU U UCUAUGCC	240	GGCAUAGA CUGAUGAG X CGAA AAAAUACA	1376
915	GUUUUUU U CUUUGCCA	241	UGGCAUAG CUGAUGAG X CGAA AAAAUAC	1377
916	UAUUUUU C UAUGCCAC	242	GUGGCAUA CUGAUGAG X CGAA AAAAAUA	1378
918	UUUUUUU A UGCCACAU	243	AUGUGGCA CUGAUGAG X CGAA AGAAAAA	1379
927	UGCCACAU U AACAUCUU	244	AAGAUGUU CUGAUGAG X CGAA AUGUGGCA	1380
928	GCCACAU A ACAUCUU	245	AAAGAUGU CUGAUGAG X CGAA AAUGUGGC	1381
933	AUUAACAU C UUUUAAAG	246	CUUUAAAA CUGAUGAG X CGAA AUGUAAU	1382
935	UAACAUU U UUAAGUU	247	AACUUUA CUGAUGAG X CGAA AGAUGUA	1383
936	AACAUCU U UAAAGUUG	248	CAACUUUA CUGAUGAG X CGAA AAGAUGU	1384
937	ACAUCUU U AAAGUUGA	249	UCAACUU CUGAUGAG X CGAA AAAGAUGU	1385
938	CAUCUUU A AAGUUGAU	250	AUCAACU CUGAUGAG X CGAA AAAAGAUG	1386
943	UUUAAAGU U GAUGAGAA	251	UUCUCAUC CUGAUGAG X CGAA ACUUUAAA	1387
953	AUGAGAAU C AAGUAUGG	252	CCAUACUU CUGAUGAG X CGAA AUUCUCAU	1388
958	AAUCAAGU A UGGAAG	253	CUUUUCCA CUGAUGAG X CGAA ACUUGAUU	1389
968	GGAAAGU A AGGCCAU	254	UAUGGCCU CUGAUGAG X CGAA ACUUUCC	1390
976	AAGGCCAU A CUCUACA	255	UGUAAGAG CUGAUGAG X CGAA AUGGCCU	1391
979	GCCAUACU C UUACUAA	256	UUAUGUAA CUGAUGAG X CGAA AGUAUGGC	1392
981	CAUACUCU U ACUAUUA	257	UAUUAUGU CUGAUGAG X CGAA AGAGUAUG	1393
982	AUACUCU A CAUAUUA	258	UUAUUAUG CUGAUGAG X CGAA AAGAGUAU	1394
986	UCUACAU A AUAAAAU	259	AAUUUAU CUGAUGAG X CGAA AUGUAAGA	1395
989	UACUAUU A AAUUCU	260	AGGAUUU CUGAUGAG X CGAA AUUAUGUA	1396
994	AAUAAAAU U CCUUUUA	261	UUAAAAGG CUGAUGAG X CGAA AUUUUAU	1397
995	AUAAAAU C CUUUUAG	262	CUUAAAAG CUGAUGAG X CGAA AAUUUAU	1398
998	AAAUCCU U UUAAGUAA	263	UUACUUA CUGAUGAG X CGAA AGGAUUU	1399
999	AAUCCUU U UAAGUAU	264	AUUACUUA CUGAUGAG X CGAA AAGGAUU	1400
1000	AUCCUUU U AAGUAUU	265	AAUACUU CUGAUGAG X CGAA AAAGGAU	1401
1001	UUCCUUU A AGUAUUU	266	AAAUACU CUGAUGAG X CGAA AAAAGGA	1402
1005	UUUUAAGU A AUUUUUC	267	GAAAAAU CUGAUGAG X CGAA ACUAAAA	1403
1008	UAAGUAU U UUUUCAA	268	UUUGAAA CUGAUGAG X CGAA AUUACUA	1404
1009	AAGUAUU U UUUCAAAG	269	CUUUGAAA CUGAUGAG X CGAA AAUACUU	1405
1010	AGUAUUU U UUCAAGA	270	UCUUUGAA CUGAUGAG X CGAA AAAUACU	1406
1011	GUAAUUU U UCAAAGAA	271	UUCUUUGA CUGAUGAG X CGAA AAAAUAC	1407
1012	UAUUUUU U CAAAGAAU	272	AUUCUUUG CUGAUGAG X CGAA AAAAAUA	1408
1013	AAUUUUU C AAAGAAUC	273	GAUUCUU CUGAUGAG X CGAA AAAAAU	1409
1021	CAAAGAAU C ACAGAAU	274	AAUUCUGU CUGAUGAG X CGAA AUUCUUUG	1410
1029	CACAGAAU U CUAGUACA	275	UGUACUAG CUGAUGAG X CGAA AUUCUGUG	1411
1030	ACAGAAU C UAGUACAU	276	AUGUACUA CUGAUGAG X CGAA AAUUCUGU	1412
1032	AGAAUUCU A GUACAUGU	277	ACAUGUAC CUGAUGAG X CGAA AGAAUUCU	1413

Table 24

1035	AUUCUAGU A CAUGUAGG	278	CCUACAUG CUGAUGAG X CGAA ACUAGAAU	1414
1041	GUACAUGU A GGUAAAUC	279	GAUUUACC CUGAUGAG X CGAA ACAUGUAC	1415
1045	AUGUAGGU A AAUCAUAA	280	UUAUGAUU CUGAUGAG X CGAA ACCUACAU	1416
1049	AGGUAAAU C AUAAAUCU	281	AGAUUUAU CUGAUGAG X CGAA AUUUACCU	1417
1052	UAAAUCAU A AAUCUGUU	282	AACAGAUU CUGAUGAG X CGAA AUGAUUUA	1418
1056	UCAUAAAU C UGUUCUAA	283	UUAGAACA CUGAUGAG X CGAA AUUUUAUGA	1419
1060	AAAUUCUGU U CUAAGACA	284	UGUCUUAG CUGAUGAG X CGAA ACAGAUUU	1420
1061	AAUCUGUU C UAAGACAU	285	AUGUCUUA CUGAUGAG X CGAA AACAGAUU	1421
1063	UCUGUUUCU A AGACAUAU	286	AUAUGUCU CUGAUGAG X CGAA AGAACAGA	1422
1070	UAAGACAU A UGAUCAAC	287	GUUGAUCA CUGAUGAG X CGAA AUGUCUUA	1423
1075	CAUAUGAU C AACAGAUG	288	CAUCUGUU CUGAUGAG X CGAA AUCAUAUG	1424
1096	CUGUGUGU U AAUAUGUG	289	CACAUUUU CUGAUGAG X CGAA ACCACCAG	1425
1097	UGGUGGUU A AUAUGUGA	290	UCACAUAU CUGAUGAG X CGAA AACCACCA	1426
1100	UGGUUAAU A UGUGACAG	291	CUGUCACA CUGAUGAG X CGAA AUUAACCA	1427
1115	AGUGAGAU U AGUCAUUA	292	AUAUGACU CUGAUGAG X CGAA AUCUCACU	1428
1116	GUGAGAUU A GUCAUAUC	293	GAUAUGAC CUGAUGAG X CGAA AAUCUCAC	1429
1119	AGAUUAGU C AUAUCACU	294	AGUGAUUA CUGAUGAG X CGAA ACUAAUCU	1430
1122	UUAGUCAU A UCACUAAU	295	AUUAGUGA CUGAUGAG X CGAA AUGACUAA	1431
1124	AGUCAUUA C ACUAAUUA	296	AUAUUAGU CUGAUGAG X CGAA AUAUGACU	1432
1128	AUAUCACU A AUUAUCUA	297	UAGUAUAU CUGAUGAG X CGAA AGUGAUUA	1433
1131	UCACUAAU A UACUAACA	298	UGUUAGUA CUGAUGAG X CGAA AUUAGUGA	1434
1133	ACUAAUUA A CUAACAAC	299	GUUGUUAG CUGAUGAG X CGAA AUUUUAGU	1435
1136	AAUAUACU A ACAACAGA	300	UCUGUUUG CUGAUGAG X CGAA AGUAUUAU	1436
1147	AACAGAAU C UAAUCUUC	301	GAAGAUUA CUGAUGAG X CGAA AUUCUGUU	1437
1149	CAGAAUCU A AUCUUCAU	302	AUGAAGAU CUGAUGAG X CGAA AGAUUCUG	1438
1152	AAUCUAAU C UUCAUUUA	303	UAAAUCAA CUGAUGAG X CGAA AUUAGAUU	1439
1154	UCUAAUCU U CAUUUAAG	304	CUUAAAUG CUGAUGAG X CGAA AGAUUAGA	1440
1155	CUAAUCUU C AUUUAAAG	305	CCUUAAAU CUGAUGAG X CGAA AAGAUUAG	1441
1158	AUCUUCAU U UAAGGCAC	306	GUGCCUUA CUGAUGAG X CGAA AUGAAGAU	1442
1159	UCUUCAUU U AAGGCACU	307	AGUGCCUU CUGAUGAG X CGAA AAUGAAGA	1443
1160	CUUCAUUU A AGGCACUG	308	CAGUGCCU CUGAUGAG X CGAA AAAUGAAG	1444
1170	GGCACUGU A GUGAAUUA	309	UAAUUCAC CUGAUGAG X CGAA ACAGUGCC	1445
1177	UAGUGAAU U AUCUGAGC	310	GCUCAGAU CUGAUGAG X CGAA AUUCACUA	1446
1178	AGUGAAUU A UCUGAGCU	311	AGCUCAGA CUGAUGAG X CGAA AAUUCACU	1447
1180	UGAAUUAU C UGAGCUAG	312	CUAGCUCA CUGAUGAG X CGAA AUAAUUCA	1448
1187	UCUGAGCU A GAGUUACC	313	GGUAACUC CUGAUGAG X CGAA AGCUCAGA	1449
1192	GCUAGAGU U ACCUAGCU	314	AGCUAGGU CUGAUGAG X CGAA ACUCUAGC	1450
1193	CUAGAGUU A CCUAGCUU	315	AAGCUAGG CUGAUGAG X CGAA AACUCUAG	1451
1197	AGUUACCU A GCUUACCA	316	UGGUAAGC CUGAUGAG X CGAA AGGUAACU	1452
1201	ACCUAGCU U ACCAUACU	317	AGUAUGGU CUGAUGAG X CGAA AGCUAGGU	1453
1202	CCUAGCUU A CCAUACUA	318	UAGUAUGG CUGAUGAG X CGAA AAGCUAGG	1454
1207	CUUACCAU A CUUAUUCU	319	AGAUAUAG CUGAUGAG X CGAA AUGGUAAG	1455
1210	ACCAUACU A UAUCUUUG	320	CAAAGAUU CUGAUGAG X CGAA AGUAUGGU	1456
1212	CAUACUAU A UCUUUGGA	321	UCCAAAGA CUGAUGAG X CGAA AUAGUAUG	1457
1214	UACUAUAU C UUUGGAAU	322	AUUCCAAU CUGAUGAG X CGAA AUUAUAGU	1458
1216	CUUAUUCU U UGGAUACA	323	UGAUUCCA CUGAUGAG X CGAA AGAUUAUG	1459
1217	UAUAUCUU U GGAAUCAU	324	AUGAUUCC CUGAUGAG X CGAA AAGAUUAU	1460

Table 24

1223	UUUGGAAU C AUGAAACC	325	GGUUUCAU CUGAUGAG X CGAA AUUCCAAA	1461
1233	UGAAACCU U AAGACUUC	326	GAAGUCUU CUGAUGAG X CGAA AGGUUUC	1462
1234	GAAACCUU A AGACUUC	327	UGAAGUCU CUGAUGAG X CGAA AAGGUUUC	1463
1240	UUAAGACU U CAGAAUGA	328	UCAUUCUG CUGAUGAG X CGAA AGUCUUA	1464
1241	UAAGACUU C AGAAUGAU	329	AUCAUUCU CUGAUGAG X CGAA AAGUCUUA	1465
1250	AGAAUGAU U UUGCAGGU	330	ACCUGCAA CUGAUGAG X CGAA AUCAUUCU	1466
1251	GAAUGAUU U UGCAGGUU	331	AACCUGCA CUGAUGAG X CGAA AAUCAUUC	1467
1252	AAUGAUUU U GCAGGUUG	332	CAACCUGC CUGAUGAG X CGAA AAAUCAUU	1468
1259	UUGCAGGU U GUCUCCA	333	UGGAAGAC CUGAUGAG X CGAA ACCUGCAA	1469
1262	CAGGUUGU C UUCAUUC	334	GAAUGGAA CUGAUGAG X CGAA ACAACCUG	1470
1264	GGUUGUCU U CCAUCCA	335	UGGA AUGG CUGAUGAG X CGAA AGACAACC	1471
1265	GUUGUCUU C CAUCCAG	336	CUGGAAUG CUGAUGAG X CGAA AAGACAAC	1472
1269	UCUCCA U. CCAGCCUA	337	UAGGCUGG CUGAUGAG X CGAA AUGGAAGA	1473
1270	CUUCCA U C CAGCCUA	338	UUAGGCUG CUGAUGAG X CGAA AAUGGAAG	1474
1277	UCCAGCCU A ACAUCCA	339	UUGGAUGU CUGAUGAG X CGAA AGGCUGGA	1475
1282	CCUAACAU C CAAUGCAG	340	CUGCAUUG CUGAUGAG X CGAA AUGUAGG	1476
1302	AGGAAAAU A AAAGAUUU	341	AAAUCUUU CUGAUGAG X CGAA AUUUUCCU	1477
1309	UAAAAGAU U UCCAGUGA	342	UCACUGGA CUGAUGAG X CGAA AUCUUUUA	1478
1310	AAAAGAUU U CCAGUGAC	343	GUCACUGG CUGAUGAG X CGAA AAUCUUUU	1479
1311	AAAGAUUU C CAGUGACA	344	UGUCACUG CUGAUGAG X CGAA AAAUCUUU	1480
1327	AGAAAAAU A UAUAUCU	345	AGAUAUA CUGAUGAG X CGAA AUUUUUCU	1481
1329	AAAAUAU A UUAUCUCA	346	UGAGAUAA CUGAUGAG X CGAA AUAUUUUU	1482
1331	AAUAUAU U AUCUCAAG	347	CUUGAGAU CUGAUGAG X CGAA AUAUAUUU	1483
1332	AAUAUAU A UCUCAAGU	348	ACUUGAGA CUGAUGAG X CGAA AAUAUAUU	1484
1334	UAUAUAU C UCAAGUAU	349	AUACUUGA CUGAUGAG X CGAA AUAUAUA	1485
1336	UAUAUAU C AAGUAUUU	350	AAAUACUU CUGAUGAG X CGAA AGAUAUA	1486
1341	UCUCAAGU A UUUUUUA	351	UUAAAAA CUGAUGAG X CGAA ACUUGAGA	1487
1343	UCAAGUAU U UUUAAAA	352	UUUUAAAA CUGAUGAG X CGAA AUACUUGA	1488
1344	CAAGUAU U UUUAAAA	353	UUUUUAAA CUGAUGAG X CGAA AAUACUUG	1489
1345	AAGUAUUU U UUUAAAA	354	AUUUUUAA CUGAUGAG X CGAA AAAUACUU	1490
1346	AGUAUUUU U UAAAAUA	355	UAUUUUUA CUGAUGAG X CGAA AAAAUACU	1491
1347	GUAUUUUU U AAAAAUA	356	AUAUUUUU CUGAUGAG X CGAA AAAAAUAC	1492
1348	UAUUUUUU A AAAAAUA	357	UAUAUUUU CUGAUGAG X CGAA AAAAAUA	1493
1354	UUAAAAAU A UAUGAAU	358	AAUUCUA CUGAUGAG X CGAA AUUUUUAA	1494
1356	AAAAUAU A UGAUUCU	359	AGAAUUC CUGAUGAG X CGAA AUAUUUUU	1495
1362	AUAUGAAU U CUCUCUCC	360	GGAGAGAG CUGAUGAG X CGAA AUUCAUAU	1496
1363	UAUGAAU C UCUCUCCA	361	UGGAGAGA CUGAUGAG X CGAA AAUCAUA	1497
1365	UGAAUUCU C UCUCCAA	362	UUUGGAGA CUGAUGAG X CGAA AGAAUUC	1498
1367	AAUUCUCU C UCCAAUA	363	UAUUUGGA CUGAUGAG X CGAA AGAGAAU	1499
1369	UUCUCUCU C CAAUAU	364	AAUAUUUG CUGAUGAG X CGAA AGAGAGAA	1500
1375	CUCCAAU A UUAACUA	365	UUAGUUA CUGAUGAG X CGAA AUUUGGAG	1501
1377	CCAAUAU U AACUAUU	366	AAUAGUU CUGAUGAG X CGAA AUAUUUGG	1502
1378	CAAAUAU A ACUAUUA	367	UAAUAGU CUGAUGAG X CGAA AAUAUUUG	1503
1382	UAUUAACU A AUUAUAG	368	CUAAUAU CUGAUGAG X CGAA AGUUAUA	1504
1385	UAACUAU U AUUAUAG	369	AAUCUAU CUGAUGAG X CGAA AUUAUUA	1505
1386	AACUAUU A UUAUAUA	370	UAAUCUA CUGAUGAG X CGAA AAUAUUA	1506
1388	CUAAUAU U AGAUAUA	371	UAUAUUCU CUGAUGAG X CGAA AUAUAUAG	1507

Table 24

1389	UAAUUAUU A GAUUAUUAU	372	AUAUAAUC CUGAUGAG X CGAA AAUAAUUA	1508
1393	UAUUAGAU U AUAUUUUG	373	CAAAAUUAU CUGAUGAG X CGAA AUCUAAUA	1509
1394	AUUAGAUU A UAUUUUGA	374	UCAAAUA CUGAUGAG X CGAA AAUCUAAU	1510
1396	UAGAUAU A UUUUGAAA	375	UUUCAAAA CUGAUGAG X CGAA AUAUUCUA	1511
1398	GAUUAUUAU U UUGAAAUG	376	CAUUUCAA CUGAUGAG X CGAA AUAUAAUC	1512
1399	AUUUAUUAU U UGAAAUGA	377	UCAUUUCA CUGAUGAG X CGAA AAUAUAAU	1513
1400	UUAUAUUU U GAAAUGAA	378	UUCAUUUC CUGAUGAG X CGAA AAAUAUAA	1514
1411	AAUGAACU U GUUGGCC	379	GGGCCAAC CUGAUGAG X CGAA AGUUCAU	1515
1414	GAACUUGU U GGCCCAUC	380	GAUGGGCC CUGAUGAG X CGAA ACAGUUC	1516
1422	UGGCCCAU C UAUUACAU	381	AUGUAAUA CUGAUGAG X CGAA AUGGGCCA	1517
1424	GCCCAUCU A UUAUACU	382	AGAUGUAA CUGAUGAG X CGAA AGAUGGGC	1518
1426	CCAUCUUAU U ACAUCUAC	383	GUAGAUGU CUGAUGAG X CGAA AUAGAUGG	1519
1427	CAUCUAUU A CAUCUACA	384	UGUAGAUG CUGAUGAG X CGAA AAUAGAUG	1520
1431	UAUUAUUAU C UACAGCUG	385	CAGCUGUA CUGAUGAG X CGAA AUGUAAUA	1521
1433	UUACAUCU A CAGCUGAC	386	GUCAGCUG CUGAUGAG X CGAA AGAUGUAA	1522
1445	CUGACCCU U GAACAUGG	387	CCAUGUUC CUGAUGAG X CGAA AGGGUCAG	1523
1458	AUGGGGGU U AGGGGAGC	388	GCUCCCCU CUGAUGAG X CGAA ACCCCCAU	1524
1459	UGGGGGUU A GGGGAGCU	389	AGCUCCCC CUGAUGAG X CGAA AACCCCA	1525
1474	CUGACAAU U CGUGGGUC	390	GACCCACG CUGAUGAG X CGAA AUUGUCAG	1526
1475	UGACAAUU C GUGGGUCC	391	GGACCCAC CUGAUGAG X CGAA AAUUGUCA	1527
1482	UCGUGGGU C CGCAAAAU	392	AUUUUGCG CUGAUGAG X CGAA ACCCACGA	1528
1491	CGCAAAAU C UUAACUAC	393	GUAGUUA CUGAUGAG X CGAA AUUUUGCG	1529
1493	CAAAUCU U AACUACCU	394	AGGUAGUU CUGAUGAG X CGAA AGAUUUUG	1530
1494	AAAUCUU A ACUACCUA	395	UAGGUAGU CUGAUGAG X CGAA AAGAUUUU	1531
1498	UCUUAACU A CCUAUAG	396	CUAUUAGG CUGAUGAG X CGAA AGUUAAGA	1532
1502	AACUACCU A AUAGCCUA	397	UAGGCUAU CUGAUGAG X CGAA AGGUAGUU	1533
1505	UACCUAAU A GCCUACUA	398	UAGUAGGC CUGAUGAG X CGAA AUUAGGUA	1534
1510	AAUAGCCU A CUAUUGAC	399	GUCAAUAG CUGAUGAG X CGAA AGGCUAUU	1535
1513	AGCCUACU A UUGACCAU	400	AUGGUCAA CUGAUGAG X CGAA AGUAGGCU	1536
1515	CCUACUUAU U GACCAUAA	401	UUAUGGUC CUGAUGAG X CGAA AUAGUAGG	1537
1522	UUGACCAU A AACCUUAC	402	GUAAGGUU CUGAUGAG X CGAA AUGGUCAA	1538
1528	AUAAACCU U ACUGAUAA	403	UUAUCAGU CUGAUGAG X CGAA AGGUUUUA	1539
1529	UAAACCUU A CUGAUAAAC	404	GUUAUCAG CUGAUGAG X CGAA AAGGUUUA	1540
1535	UUACUGAU A ACAUAAAC	405	GUUUUAGU CUGAUGAG X CGAA AUCAGUAA	1541
1540	GAUAAACU A AACAGUAA	406	UUACUGUU CUGAUGAG X CGAA AUGUUAUC	1542
1547	UAAACAGU A AAUUAACA	407	UGUUAAUU CUGAUGAG X CGAA ACUGUUUA	1543
1551	CAGUAAAU U AACACUA	408	UAUGUGUU CUGAUGAG X CGAA AUUUACUG	1544
1552	AGUAAAUU A ACACUAU	409	AUAUGUGU CUGAUGAG X CGAA AAUUUACU	1545
1559	UAACACAU A UUUUGCGU	410	ACGCAAAA CUGAUGAG X CGAA AUGUGUUA	1546
1561	ACACUAU U UUGCGUGU	411	ACACGCAA CUGAUGAG X CGAA AUAUGUGU	1547
1562	CACAUAAU U UGCGUGUU	412	AACACGCA CUGAUGAG X CGAA AAUAUGUG	1548
1563	ACAUAAUU U GCGUGUUA	413	UAACACGC CUGAUGAG X CGAA AAAUAUGU	1549
1570	UUGCGUGU U AUAUGUAU	414	AUACAUAU CUGAUGAG X CGAA ACACGCAA	1550
1571	UGCGUGUU A UAUGUAU	415	AAUACAUA CUGAUGAG X CGAA AACACGCA	1551
1573	CGUGUUAU A UGUUAUUA	416	AUAUAACA CUGAUGAG X CGAA AUAACACG	1552
1577	UUAUAUGU A UUAUACAC	417	GUGUAUAA CUGAUGAG X CGAA ACAUAUAA	1553
1579	AUAUGUAU U AUACACUA	418	UAGUGUAU CUGAUGAG X CGAA AUACAUAU	1554

Table 24

1580	UAUGUAUU A UACACUAU	419	AUAGUGUA CUGAUGAG X CGAA AAUACAU	1555
1582	UGUAUUUU A CACUAUUU	420	AUAUAGUG CUGAUGAG X CGAA AUAAUACA	1556
1587	UAUACACU A UAUUCCUA	421	UAGGAAUA CUGAUGAG X CGAA AGUGUAUA	1557
1589	UACACUAU A UUCCUACA	422	UGUAGGAA CUGAUGAG X CGAA AUAGUGUA	1558
1591	CACUAUUU U CCUACAAU	423	AUUGUAGG CUGAUGAG X CGAA AUUAGUG	1559
1592	ACUAUAUU C CUACAAUA	424	UAUUGUAG CUGAUGAG X CGAA AAUAUAGU	1560
1595	AUAUCCU A CAUAAAAG	425	CUUUUUG CUGAUGAG X CGAA AGGAAUAU	1561
1600	CCUACAAU A AAGUAAGC	426	GCUUACUU CUGAUGAG X CGAA AUUGUAGG	1562
1605	AAUAAAAGU A AGCUAGAG	427	CUCUAGCU CUGAUGAG X CGAA ACUUUAUU	1563
1610	AGUAAGCU A GAGAAAUA	428	AUUUUCUC CUGAUGAG X CGAA AGCUUACU	1564
1621	GAAAAGU U AUUUAGAA	429	UUCUAAA CUGAUGAG X CGAA ACAUUUUC	1565
1622	AAAAUGUU A UUUAGAAA	430	UUUCUAAA CUGAUGAG X CGAA AACAUUUU	1566
1624	AAUGUAUU U UAGAAAUA	431	AUUUUCUA CUGAUGAG X CGAA AUAACAUA	1567
1625	AUGUAUUU U AGAAAUAU	432	GAUUUUCU CUGAUGAG X CGAA AAUAACAU	1568
1626	UGUAUUUU A GAAAUAU	433	UGAUUUUC CUGAUGAG X CGAA AAAUAACA	1569

Input Sequence = PLN. Cut Site = UH/.

Stem Length = 8. Core Sequence = CUGAUGAG X CGAA (X = GCCGUUAGGC or other stem II)

PLN (Homo sapiens phospholamban (PLN) mRNA.; 1635 bp)

Table 25

Table 25: Human Phospholamban (PLN) NCH Ribozyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
15	CAGAAAAC U AGCUAAAC	434	GUUUAGCU CUGAUGAG X CGAA IUUUUCUG	1570
17	GAAACUC C CUAAACAC	435	GUGUUUAG CUGAUGAG X CGAA IAGUUUUC	1571
18	AAACUCC C UAAACACC	436	GGUGUUUA CUGAUGAG X CGAA IGAGUUUU	1572
19	AAACUCC C AACACCC	437	GGGUGUUU CUGAUGAG X CGAA IGGAGUUU	1573
20	AACUCCC A AACACCCG	438	CGGGUGUU CUGAUGAG X CGAA IGGGAGUU	1574
23	UCCCCAGC U ACCCGUAA	439	UUACGGGU CUGAUGAG X CGAA ICUGGGGA	1575
28	AGCUAAAC A UAAGACUU	440	AAGUCUUA CUGAUGAG X CGAA IUUUAGCU	1576
30	CUAAACAC C AGACUUA	441	UGAAGUCU CUGAUGAG X CGAA IUGUUUAG	1577
31	UAAACACC C GACUUAU	442	AUGAAGUC CUGAUGAG X CGAA IGUGUUUA	1578
39	CGUAAGAC U ACAACACA	443	UGUGUUGU CUGAUGAG X CGAA IUCUUAACG	1579
42	AAGACUUC A ACACAAUA	444	UAUUGUGU CUGAUGAG X CGAA IAAGUCUU	1580
46	CUUCAUAC A AAUACUCU	445	AGAGUAUU CUGAUGAG X CGAA IUAUGAAG	1581
49	CAUACAAC A ACUCUAUA	446	UAUAGAGU CUGAUGAG X CGAA IUUGUAUG	1582
51	UACAACAC A UCUAUACU	447	AGUAUAGA CUGAUGAG X CGAA IUGUUGUA	1583
56	CACAAUAC U ACUGUGAU	448	AUCACAGU CUGAUGAG X CGAA IUAUUGUG	1584
58	CAAUACUC U UGUGAUGA	449	UCAUCACA CUGAUGAG X CGAA IAGUAUUG	1585
63	CUCUAUAC U UGAUCACA	450	UGUGAUCA CUGAUGAG X CGAA IUAUAGAG	1586
73	UGAUGAUC A UGCCAAGG	451	CCUUGGCA CUGAUGAG X CGAA IAUCAUCA	1587
75	AUGAUCAC A CCAAGGCU	452	AGCCUUGG CUGAUGAG X CGAA IUGAUCAU	1588
78	AUCACAGC U AGGCUACC	453	GGUAGCCU CUGAUGAG X CGAA ICUGUGAU	1589
81	ACAGCUGC C CUACCUAA	454	UUAGGUAG CUGAUGAG X CGAA ICAGCUGU	1590
82	CAGCUGCC A UACCUAAA	455	UUUAGGUA CUGAUGAG X CGAA IGCAGCUG	1591
87	GCCAAGGC U AAAAGAAG	456	CUUCUUUU CUGAUGAG X CGAA ICCUUGGC	1592
90	AAGGCUAC C AGAAGACA	457	UGUCUUCU CUGAUGAG X CGAA IUAGCCUU	1593
91	AGGCUACC U GAAGACAG	458	CUGUCUUC CUGAUGAG X CGAA IGUAGCCU	1594
102	AAGAAGAC A UCUCUAU	459	AUAUGAGA CUGAUGAG X CGAA IUCUUCUU	1595
109	CAGUUAUC U UUUGGCUG	460	CAGCCAAA CUGAUGAG X CGAA IAUAAACUG	1596
111	GUUAUCUC A UGGCUGCC	461	GGCAGCCA CUGAUGAG X CGAA IAGUAAC	1597
120	UAUUUGGC U GCUUUUUA	462	UAAAAGC CUGAUGAG X CGAA ICCAAUA	1598
123	UUGGCUGC C UUUUAUCU	463	AGAUAAA CUGAUGAG X CGAA ICAGCCAA	1599
124	UGGCUGCC A UUUUAUCU	464	AAGAUAAA CUGAUGAG X CGAA IGCAGCCA	1600
127	CUGCCAGC U AUCUUUCU	465	AGAAAGAU CUGAUGAG X CGAA ICUGGCAG	1601
135	UUUUUAUC U CUCGACCA	466	UGGUCGAG CUGAUGAG X CGAA IAUAAAAA	1602
139	UAUCUUUC U ACCACUUA	467	UAAGUGGU CUGAUGAG X CGAA IAAAGAU	1603
141	UCUUUCUC U CACUAAA	468	UUUAAGUG CUGAUGAG X CGAA IAGAAAGA	1604
146	CUCUCGAC C AAAACUUC	469	GAAGUUUU CUGAUGAG X CGAA IUCGAGAG	1605
147	UCUCGACC A AAACUUA	470	UGAAGUUU CUGAUGAG X CGAA IGUCGAGA	1606
149	UCGACCAC U ACUUCAGA	471	UCUGAAGU CUGAUGAG X CGAA IUGGUCGA	1607
156	CUUAAAAC U ACUCCUG	472	CAGGAAGU CUGAUGAG X CGAA IUUUUAAG	1608
159	AAAACUUC A UCCUGUCC	473	GGACAGGA CUGAUGAG X CGAA IAAGUUUU	1609
163	CUUCAGAC U GUCCUGCU	474	AGCAGGAC CUGAUGAG X CGAA IUCUGAAG	1610
166	CAGACUUC C CUGCUGGU	475	ACCAGCAG CUGAUGAG X CGAA IAAGUCUG	1611
167	AGACUUC U UGCUGGUA	476	UACCAGCA CUGAUGAG X CGAA IGAAGUCU	1612
171	UUCUGUC C GGUAUCAU	477	AUGAUACC CUGAUGAG X CGAA IACAGGAA	1613

Table 25

172	UCCUGUCC U GUAUCAUG	478	CAUGAUAC CUGAUGAG X CGAA IGACAGGA	1614
175	UGUCCUGC U UCAUGGAG	479	CUCCAUGA CUGAUGAG X CGAA ICAGGACA	1615
182	CUGGUAUC A GAAAGUCC	480	GGACUUUC CUGAUGAG X CGAA IAUACCAG	1616
194	AGAAAGUC C CCUCACUC	481	GAGUGAGG CUGAUGAG X CGAA IACUUUCU	1617
195	GAAAGUCC A CUCACUCG	482	CGAGUGAG CUGAUGAG X CGAA IGACUUUC	1618
200	UCCAUAAC C UCGCUCAG	483	CUGAGCGA CUGAUGAG X CGAA IUAUUGGA	1619
201	CCAAUACC U CGCUCAGC	484	GCUGAGCG CUGAUGAG X CGAA IGUAUUGG	1620
203	AAUACCUC A CUCAGCUA	485	UAGCUGAG CUGAUGAG X CGAA IAGGUUUU	1621
205	UACCUCAC U CAGCUAUA	486	UAUAGCUG CUGAUGAG X CGAA IUGAGGUA	1622
209	UCACUCGC U UAUAAGAA	487	UUCUUAUA CUGAUGAG X CGAA ICAGUGA	1623
211	ACUCGCUC A UAAGAAGA	488	UCUUCUUA CUGAUGAG X CGAA IAGCGAGU	1624
214	CGCUCAGC U GAAGAGCC	489	GGCUCUUC CUGAUGAG X CGAA ICUGAGCG	1625
226	AGAAGAGC C CCAUUGAA	490	UUCAAUGG CUGAUGAG X CGAA ICUCUUCU	1626
227	GAAGAGCC U CAUUGAAA	491	UUUCAAUG CUGAUGAG X CGAA IGCUCUUC	1627
229	AGAGCCUC A UUGAAUUG	492	CAUUUCAA CUGAUGAG X CGAA IAGGCUCU	1628
232	GCCUCAAC C AAAUGCCU	493	AGGCAUUU CUGAUGAG X CGAA IUUGAGGC	1629
233	CCUCAACC A AAUGCCUC	494	GAGGCAUU CUGAUGAG X CGAA IGUUGAGG	1630
243	UGAAAUUG C CAAGCACG	495	CGUGCUUG CUGAUGAG X CGAA ICAUUUCA	1631
244	GAAAUGCC U AAGCACGU	496	ACGUGCUU CUGAUGAG X CGAA IGCAUUUC	1632
246	AAUGCCUC A GCACGUCA	497	UGACGUGC CUGAUGAG X CGAA IAGGCAUU	1633
249	GCCUCAAC A CGUCAAAA	498	UUUUGACG CUGAUGAG X CGAA IUUGAGGC	1634
253	CAACAAGC A AAAAGCUA	499	UAGCUUUU CUGAUGAG X CGAA ICUGUUUG	1635
258	AGCACGUC A CUACAGAA	500	UUCUGUAG CUGAUGAG X CGAA IACGUGCU	1636
264	UCAAAAGC U AAUCUAUU	501	AAUAGAUU CUGAUGAG X CGAA ICUUUUGA	1637
267	AAAGCUAC A CUAUUUAU	502	AUAAAUAG CUGAUGAG X CGAA IUAGCUUU	1638
273	ACAGAAUC U AUCAAUUU	503	AAAUUGAU CUGAUGAG X CGAA IAUUCUGU	1639
281	UAUUUAUC A CUGUCUCA	504	UGAGACAG CUGAUGAG X CGAA IAUAAUA	1640
287	UCAAUUUC U CAUCUUA	505	UUAAGAUG CUGAUGAG X CGAA IAAAUUGA	1641
291	UUUCUGUC U UUAUAUUG	506	CAUAUUAA CUGAUGAG X CGAA IACAGAAA	1642
293	UCUGUCUC A AAUAUGUC	507	GACAUUUU CUGAUGAG X CGAA IAGACAGA	1643
296	GUCUCAUC U AUGUCUCU	508	AGAGACAU CUGAUGAG X CGAA IAUGAGAC	1644
306	AAUAUGUC U CUGAUCUG	509	CAGAUCAG CUGAUGAG X CGAA IACAUUUU	1645
308	UAUGUCUC U GAUCUGUA	510	UACAGAUU CUGAUGAG X CGAA IAGACAUU	1646
312	UCUCUUGC U UGUUAUUA	511	AUGAUACA CUGAUGAG X CGAA ICAAGAGA	1647
317	UGCUGAUC U CAUCGUGA	512	UCACGAUG CUGAUGAG X CGAA IAUACAGA	1648
323	UCUGUAUC A GAUCUUC	513	GAAGCAUC CUGAUGAG X CGAA IAUACAGA	1649
333	CGUGAUGC U UGAAGUUC	514	GAACUUCA CUGAUGAG X CGAA ICAUCACG	1650
336	GAUGCUUC U AGUUCUGC	515	GCAGAAUC CUGAUGAG X CGAA IAAGCAUC	1651
338	UGCUCUCUC U UUCUGCUA	516	UAGCAGAA CUGAUGAG X CGAA IAGAAGCA	1652
346	UGAAGUUC U CAACCUCU	517	AGAGGUUG CUGAUGAG X CGAA IAACUUCA	1653
349	AGUUCUGC U CCUCUAGA	518	UCUAGAGG CUGAUGAG X CGAA ICAGAACU	1654
352	UCUGCUAC A CUAGAUCU	519	AGAUCUAG CUGAUGAG X CGAA IUAGCAGA	1655
355	GCUACAAC C GAUCUGCA	520	UGCAGAUU CUGAUGAG X CGAA IUUGUAGC	1656
356	CUACAACC U AUCUGCAG	521	CUGCAGAU CUGAUGAG X CGAA IGUUGUAG	1657
358	ACAACCUC U CUGCAGCU	522	AGCUGCAG CUGAUGAG X CGAA IAGGUUGU	1658
364	UCUAGAUC U CUUGCCAC	523	GUGGCAAG CUGAUGAG X CGAA IAUUCAGA	1659
367	AGAUCUGC A GCCACAUC	524	GAUGUGGC CUGAUGAG X CGAA ICAGAUCU	1660

Table 25

370	UCUGCAGC U ACAUCAGC	525	GCUGAUGU CUGAUGAG X CGAA ICUGCAGA	1661
374	CAGCUUGC C CAGCUUAA	526	UUAAGCUG CUGAUGAG X CGAA ICAAGCUG	1662
375	AGCUUGCC A AGCUUAAA	527	UUUAAGCU CUGAUGAG X CGAA IGCAAGCU	1663
377	CUUGCCAC A CUUAAAAU	528	AUUUUAAG CUGAUGAG X CGAA IUGGCAAG	1664
380	GCCACAUC A AAAAUCUG	529	CAGAUUUU CUGAUGAG X CGAA IAUGUGGC	1665
383	ACAUCAGC U AUCUGUCA	530	UGACAGAU CUGAUGAG X CGAA ICUGAUGU	1666
391	UUA AAAAUC U UCCCAUGC	531	GCAUGGGA CUGAUGAG X CGAA IAUUUUAA	1667
395	AAUCUGUC A AUGCAGAC	532	GUCUGCAU CUGAUGAG X CGAA IACAGAUU	1668
398	CUGUCAUC C CAGACAGG	533	CCUGUCUG CUGAUGAG X CGAA IAUGACAG	1669
399	UGUCAUCC C AGACAGGA	534	UCCUGUCU CUGAUGAG X CGAA IGAUGACA	1670
400	GUCAUCCC A GACAGGAA	535	UUCUGUC CUGAUGAG X CGAA IGGAUGAC	1671
404	UCCCAUGC A GGAAACA	536	UGUUUCC CUGAUGAG X CGAA ICAUGGGA	1672
408	AUGCAGAC A AACAAUUAU	537	AUAUUGUU CUGAUGAG X CGAA IUCUGCAU	1673
416	AGGAAAAC A UGUUAUAC	538	GUUAUACA CUGAUGAG X CGAA IUUUUCCU	1674
429	UGUAUAAC A ACUUCUG	539	CAGGAAGU CUGAUGAG X CGAA IUUAUACA	1675
433	UAACAGAC C CCUGAGUA	540	UACUCAGG CUGAUGAG X CGAA IUCUGUUA	1676
434	AACAGACC A CUGAGUAG	541	CUACUCAG CUGAUGAG X CGAA IGUCUGUU	1677
436	CAGACCAC U GAGUAGAA	542	UUCUACUC CUGAUGAG X CGAA IUGGUCUG	1678
439	ACCACUUC C UAGAAGAG	543	CUCUUCUA CUGAUGAG X CGAA IAAGUGGU	1679
440	CCACUUC U AGAAGAGU	544	ACUCUUCU CUGAUGAG X CGAA IGAAGUGG	1680
456	AGAGUUUC U GAAAAGGU	545	ACCUUUUC CUGAUGAG X CGAA IAAACUCU	1681
470	AAAAGGUC A UAAGACUA	546	UAGUCUUA CUGAUGAG X CGAA IACCUUUU	1682
481	AUUAAAGAC U CUUAUUGU	547	ACAAUAAG CUGAUGAG X CGAA IUCUUAU	1683
487	ACUAAAAC U GUUACCAU	548	AUGGUAAC CUGAUGAG X CGAA IUUUUAGU	1684
497	AUUGUUAC C GUUUCAU	549	AUGAAUAC CUGAUGAG X CGAA IUAACAAU	1685
498	UUGUUACC A UAUUCAUC	550	GAUGAAUA CUGAUGAG X CGAA IGUAACAA	1686
508	AUGUAUUC A UUGGAUCU	551	AGAUCCAA CUGAUGAG X CGAA IAAUACAU	1687
511	UAUUCAUC U GAUCUUGU	552	ACAAGAUC CUGAUGAG X CGAA IAUGAAUA	1688
520	GUUGGAUC U AACAUCAA	553	UUCAUGUU CUGAUGAG X CGAA IAUCCAAC	1689
528	UUGUAAAC A AAGGGCUU	554	AAGCCCUU CUGAUGAG X CGAA IUUUACAA	1690
539	AAAAGGGC U UUUCAAAA	555	UUUUGAAA CUGAUGAG X CGAA ICCCUUUU	1691
548	UUUUUUUC A UUAACUUC	556	GAAGUUAA CUGAUGAG X CGAA IAAAUAUA	1692
558	AAAUUAAC U AAUAAGUG	557	CACUUAUU CUGAUGAG X CGAA IUUAUUUU	1693
561	UUAACUUC A AAGUGUAU	558	AUACACUU CUGAUGAG X CGAA IAAGUUAA	1694
581	UAAAAGUC A UUGAUUUC	559	GAAAUCAA CUGAUGAG X CGAA ICAUUUUA	1695
584	AAUGCAAC U AUUCCUC	560	GAGGAAAU CUGAUGAG X CGAA IUUGCAUU	1696
594	UUGAUUUC C CAUGGCUC	561	GAGCCAUG CUGAUGAG X CGAA IAAAUCAA	1697
595	UGAUUUCC U AUGGCUCA	562	UGAGCAU CUGAUGAG X CGAA IGAAAUCA	1698
597	AUUUCCUC A GGCUCACA	563	UGUGAGCC CUGAUGAG X CGAA IAGGAAAU	1699
600	UCCUCAAC A UCACAAAU	564	AUUUGUGA CUGAUGAG X CGAA IUUGAGGA	1700
605	AACAUGGC U AAUUUCUA	565	UAGAAAUU CUGAUGAG X CGAA ICCAUGUU	1701
607	CAUGGCUC A UUUUAUC	566	GAUAGAAA CUGAUGAG X CGAA IAGCCAUG	1702
609	UGGCUCAC A UCUAUCCC	567	GGGAUAGA CUGAUGAG X CGAA IUGAGCCA	1703
616	CAAAUUUC U CAAAUUUU	568	AAGAUUUG CUGAUGAG X CGAA IAAAUUUG	1704
620	UUUCUAUC C UCUUUUCU	569	AGAAAAGA CUGAUGAG X CGAA IAUAGAAA	1705
621	UUCUAUCC C CUUUUCUG	570	CAGAAAAG CUGAUGAG X CGAA IGAUAGAA	1706
622	UCUAUCCC A UUUUCUGA	571	UCAGAAAA CUGAUGAG X CGAA IGGAUAGA	1707

Table 25

627	CCCCAAUC U UGAAGAUG	572	CAUCUUCA CUGAUGAG X CGAA IAUUUGGG	1708
632	AUCUUUUC U AUGAAGAG	573	CUCUUCAU CUGAUGAG X CGAA IAAAAGAU	1709
659	UUUAAAAC U UGCCAACA	574	UGUUGGCA CUGAUGAG X CGAA IUUUUAAA	1710
662	AAAACUGC A CAACAAGU	575	ACUUGUUG CUGAUGAG X CGAA ICAGUUUU	1711
664	AACUGCAC U ACAAGUUC	576	GAACUUGU CUGAUGAG X CGAA IUGCAGUU	1712
667	UGCACUGC C AGUUCACU	577	AGUGAACU CUGAUGAG X CGAA ICAGUGCA	1713
668	GCACUGCC A GUUCACUU	578	AAGUGAAC CUGAUGAG X CGAA IGCAGUGC	1714
671	CUGCCAAC A CACUUCAU	579	AUGAAGUG CUGAUGAG X CGAA IUUGGCAG	1715
677	ACAAGUUC A AUAUAUAA	580	UUAUAUUA CUGAUGAG X CGAA IAACUUGU	1716
679	AAGUUCAC U AUAUAAAG	581	CUUUAUUA CUGAUGAG X CGAA IUGAACUU	1717
682	UUCACUUC A UAAAGCAU	582	AUGCUUUA CUGAUGAG X CGAA IAAGUGAA	1718
693	UAUAAAGC A UUUUACUC	583	GAGUAAAA CUGAUGAG X CGAA ICUUUAUA	1719
704	AUUUUUAC U UGAGGUGA	584	UCACCUCA CUGAUGAG X CGAA IUAAAAAU	1720
706	UUUUACUC U AGGUGAAU	585	AUUCACCU CUGAUGAG X CGAA IAGUAAAA	1721
733	UAUAUUAC A AAAAGCUU	586	AAGCUUUU CUGAUGAG X CGAA IUAAUAUA	1722
744	GUAAAAGC U UAAUACUA	587	UAGUAUUA CUGAUGAG X CGAA ICUUUUAC	1723
747	AAAGCUUC U UACUAAGU	588	ACUUAUGA CUGAUGAG X CGAA IAAGCUUU	1724
755	UUUAAUAC U AUUUUUCA	589	UGAAAAAU CUGAUGAG X CGAA IUAUUAAA	1725
767	UAUUUUUC A UUCACCAA	590	UUGGUGAA CUGAUGAG X CGAA IAAAAUAU	1726
772	UUCAGGUC U CAAGUAUC	591	GAUACUUG CUGAUGAG X CGAA IACCUGAA	1727
775	AGGUCUUC A GUAUCAA	592	UUUGAUAC CUGAUGAG X CGAA IAAGACCU	1728
777	GUCUUCAC C AUCAAAGU	593	ACUUUGAU CUGAUGAG X CGAA IUGAAGAC	1729
778	UCUUCACC A UCAAAGUA	594	UACUUUGA CUGAUGAG X CGAA IGUGAAGA	1730
785	CAAGUAUC A AAUACAC	595	GUGUUAAU CUGAUGAG X CGAA IAUACUUG	1731
796	GUAAUAAC A UGAAGUGU	596	ACACUUCA CUGAUGAG X CGAA IUUAUUAC	1732
798	AAUAACAC A AAGUGUCA	597	UGACACUU CUGAUGAG X CGAA IUGUUAUU	1733
810	GAAGUGUC A UCAAAUA	598	UAUUUUGA CUGAUGAG X CGAA IACACUUC	1734
817	CAUUAUUC A AGUCCACU	599	AGUGGACU CUGAUGAG X CGAA IAAUAAUG	1735
826	AAUAGUC C ACUCCUCA	600	UGAGGAGU CUGAUGAG X CGAA IACUAUUU	1736
827	AAUAGUCC A CUCCUCAC	601	GUGAGGAG CUGAUGAG X CGAA IGACUAUU	1737
829	UAGUCCAC U CCUCACAU	602	AUGUGAGG CUGAUGAG X CGAA IUGGACUA	1738
833	CCACUGAC U ACAUCUGU	603	ACAGAUGU CUGAUGAG X CGAA IUCAGUGG	1739
835	ACUGACUC C AUCUGUUA	604	UAACAGAU CUGAUGAG X CGAA IAGUCAGU	1740
836	CUGACUCC U UCUGUUAU	605	AUAACAGA CUGAUGAG X CGAA IGAGUCAG	1741
838	GACUCCUC A UGUUAUCU	606	AGAUAAAC CUGAUGAG X CGAA IAGGAGUC	1742
840	CUCCUCAC A UUAUCUUA	607	UAAGAUAA CUGAUGAG X CGAA IUGAGGAG	1743
843	CUCACAUC U UCUUAUUA	608	UAAUAAGA CUGAUGAG X CGAA IAUGUGAG	1744
850	CUGUUAUC U AUAAGAA	609	UUCUUUAU CUGAUGAG X CGAA IAUAAACAG	1745
864	UAAAGAAC U GUAGUAAC	610	GUUACUAC CUGAUGAG X CGAA IUUCUUUA	1746
877	GUAGUAAC U GAAUCUAC	611	GUAGAUUC CUGAUGAG X CGAA IUUACUAC	1747
881	UAACUAUC A CUACAUC	612	GAAUGUAG CUGAUGAG X CGAA IAUAGUUA	1748
887	UCAGAAUC U UCUAAAAC	613	GUUUUAGA CUGAUGAG X CGAA IAUUCUGA	1749
890	GAAUCUAC A AAAACAGA	614	UCUGUUUU CUGAUGAG X CGAA IUAGAUC	1750
894	CUACAUC U CAGAAAUU	615	AAUUUCUG CUGAUGAG X CGAA IAAUGUAG	1751
900	UCUAAAAC A UUGUAUUU	616	AAAUACAA CUGAUGAG X CGAA IUUUUAGA	1752
917	AUUUUUUC U CACAUUAA	617	UUAAUGUG CUGAUGAG X CGAA IAAAAAU	1753
922	UUCUAUGC C UAACAUCU	618	AGAUGUUA CUGAUGAG X CGAA ICAUAGAA	1754

Table 25

923	UCUAUGCC A AACAUUU	619	AAGAUGUU CUGAUGAG X CGAA IGCAUAGA	1755
925	UAUGCCAC A CAUCUUU	620	AAAAGAUG CUGAUGAG X CGAA IUGGCAUA	1756
931	ACAUUAA C A UUAAGUU	621	AACUUUAA CUGAUGAG X CGAA IUUAAUGU	1757
934	UUAACAUC U AAGUUGAU	622	AUCAACUU CUGAUGAG X CGAA IAUGUUA	1758
954	UGAGAAUC A UGGAAAAG	623	CUUUUCCA CUGAUGAG X CGAA IAUUCUCA	1759
973	AGUAAGGC C UCUAACAU	624	AUGUAAGA CUGAUGAG X CGAA ICCUUAU	1760
974	GUAAGGCC A CUUACUA	625	UAUGUAAG CUGAUGAG X CGAA IGCCUUA	1761
978	GGCCAUAC U CAUAAUA	626	UUUAUAUG CUGAUGAG X CGAA IUAUGGCC	1762
980	CCAUACUC U UAAUAAA	627	UUUUUUA CUGAUGAG X CGAA IAGUAUGG	1763
984	ACUCUUA C A AAAAUUC	628	GGAAUUU CUGAUGAG X CGAA IUAAGAGU	1764
996	UAAAAUUC C AAGUAUU	629	AAUUACUU CUGAUGAG X CGAA IAAUUUUA	1765
997	AAAAUUC U AGUAAUU	630	AAAUUACU CUGAUGAG X CGAA IGAAUUU	1766
1014	AUUUUUUC A AUCACAGA	631	UCUGUGAU CUGAUGAG X CGAA IAAAAAU	1767
1022	AAAGAAUC A AUUCUAGU	632	ACUAGAAU CUGAUGAG X CGAA IAUUCUU	1768
1024	AGAAUCAC A UCUGUAC	633	GUACUAGA CUGAUGAG X CGAA IUGAUUCU	1769
1031	CAGAAUUC U CAUGUAGG	634	CCUACAUG CUGAUGAG X CGAA IAAUUCUG	1770
1037	UCUAGUAC A GGUAUUC	635	GAUUUACC CUGAUGAG X CGAA IUACUAGA	1771
1050	GGUAAUUC A UCUGUUCU	636	AGAACAGA CUGAUGAG X CGAA IAUUUACC	1772
1057	CAUAAUUC U UAAGACAU	637	AUGUCUUA CUGAUGAG X CGAA IAUUUUAG	1773
1062	AUCUGUUC U CAUAUGAU	638	AUCAUAUG CUGAUGAG X CGAA IAAACAGU	1774
1068	UCUAAGAC A AUCAACAG	639	CUGUUGAU CUGAUGAG X CGAA IUCUAGA	1775
1076	AUAUGAUC A AUGAGAAC	640	GUUCUCAU CUGAUGAG X CGAA IAUCAUAU	1776
1079	UGAUCAAC A AGAACUGG	641	CCAGUUCU CUGAUGAG X CGAA IUUGAUCA	1777
1089	AUGAGAAC U GUUAUAU	642	AUAUUAAC CUGAUGAG X CGAA IUUCUCAU	1778
1107	UAUGUGAC A GAUUAGUC	643	GACUAAUC CUGAUGAG X CGAA IUCACUA	1779
1120	GAUUAGUC A ACUAAUAU	644	AUAUUAGU CUGAUGAG X CGAA IACUAAUC	1780
1125	GUCAUAUC A UAUACUA	645	UUAGUAUA CUGAUGAG X CGAA IAUUAUGAC	1781
1127	CAUAUCAC U UACUAACA	646	UGUUAGUA CUGAUGAG X CGAA IUGUAUUG	1782
1135	UAAUAUAC U ACAGAAUC	647	GAUUCUGU CUGAUGAG X CGAA IUAUAUUA	1783
1139	AUACUAAC A AAUCUAAU	648	AUUAGAUU CUGAUGAG X CGAA IUUAGUAU	1784
1142	CUACAAC A CUAAUCUU	649	AAGAUUAG CUGAUGAG X CGAA IUUGUUAG	1785
1148	ACAGAAUC U UUCAUUUA	650	UAAAUGAA CUGAUGAG X CGAA IAUUCUGU	1786
1153	AUCUAAUC U UUAAGGCA	651	UGCCUUA CUGAUGAG X CGAA IAUUAGAU	1787
1156	UAAUCUUC A AGGCACUG	652	CAGUGCCU CUGAUGAG X CGAA IAAAGUUA	1788
1165	UUUAAGGC A AGUGAAUU	653	AAUUCACU CUGAUGAG X CGAA ICCUAAA	1789
1167	UAAGGCAC U UGAUUUAU	654	AUAUUUCA CUGAUGAG X CGAA IUGCCUUA	1790
1181	GAAUUAUC U UAGAGUUA	655	UAACUCUA CUGAUGAG X CGAA IAUAAUUC	1791
1186	AUCUGAGC U UUACCUAG	656	CUAGGUAA CUGAUGAG X CGAA ICUCAGAU	1792
1195	AGAGUUA C UUAACUA	657	UAUGGUAA CUGAUGAG X CGAA IUAACUCU	1793
1196	GAGUUAAC U UACCAUAC	658	GUAUGGUA CUGAUGAG X CGAA IGUAACUC	1794
1200	UACCUAGC U AUACUAUA	659	UAUAGUAU CUGAUGAG X CGAA ICUAGGUA	1795
1204	UAGCUUAC C UAUUCUU	660	AAGAUUA CUGAUGAG X CGAA IUAAGCUA	1796
1205	AGCUUACC A AUAUCUU	661	AAAGAUUA CUGAUGAG X CGAA IGUAAGCU	1797
1209	UACCAUAC U CUUUGGAA	662	UUCCAAAG CUGAUGAG X CGAA IUAUGGUA	1798
1215	ACUAUAUC U AAUCAUGA	663	UCAUGAUU CUGAUGAG X CGAA IAUUAUGU	1799
1224	UUGGAAUC A ACCUUAAG	664	CUUAAGGU CUGAUGAG X CGAA IAUUCCAA	1800
1231	CAUGAAAC C GACUUCAG	665	CUGAAGUC CUGAUGAG X CGAA IUUUCUAG	1801

Table 25

1232	AUGAAACC U ACUUCAGA	666	UCUGAAGU CUGAUGAG X CGAA IGUUUCAU	1802
1239	CUUAAGAC U AAUGAUUU	667	AAAUCAUU CUGAUGAG X CGAA IUCUUAAG	1803
1242	AAGACUUC A GAUUUUGC	668	GCAAAAUC CUGAUGAG X CGAA IAAGUCUU	1804
1255	GAUUUUGC A GUCUCCA	669	UGGAAGAC CUGAUGAG X CGAA ICAAAAUC	1805
1263	AGGUUGUC U UCCAGCC	670	GGCUGGAA CUGAUGAG X CGAA IACAACCU	1806
1266	UUGUCUUC C CAGCCUAA	671	UUAGGCUG CUGAUGAG X CGAA IAAGACAA	1807
1267	UGUCUUC C AAGCCUAA	672	GUUAGGCU CUGAUGAG X CGAA IGAAGACA	1808
1271	UUCCAUC C UAACAUC	673	GGAUGUUA CUGAUGAG X CGAA IAAUGGAA	1809
1272	UCCAUC C AACAUCCA	674	UGGAUGUU CUGAUGAG X CGAA IGAAUGGA	1810
1275	AUUCAGC C AUCCAUG	675	CAUUGGAU CUGAUGAG X CGAA ICUGGAAU	1811
1276	UUCAGCC U UCCAUGC	676	GCAUUGGA CUGAUGAG X CGAA IGCUGGAA	1812
1280	AGCCUAA C AUGCAGGC	677	GCCUGCAU CUGAUGAG X CGAA IUUAGGCU	1813
1283	CUAACAUC C CAGGCAAG	678	CUUGCCUG CUGAUGAG X CGAA IAUGUAG	1814
1284	UAACAUC C AGGCAAG	679	CCUUGCCU CUGAUGAG X CGAA IGAUGUA	1815
1289	UCCAUGC A AGGAAAU	680	AUUUCCU CUGAUGAG X CGAA ICAUUGGA	1816
1293	AUGCAGC A AAUAAAA	681	UUUUUUU CUGAUGAG X CGAA ICCUGCAU	1817
1312	AAGAUUC C ACAGAAA	682	UUUUCUGU CUGAUGAG X CGAA IAAAUUU	1818
1313	AGAUUUC A CAGAAAA	683	UUUUCUG CUGAUGAG X CGAA IGAAAUUU	1819
1319	CCAGUGAC A AAUAUAU	684	AAUAUAU CUGAUGAG X CGAA IUCACUGG	1820
1335	AUAUAUC U UAUAUAU	685	AAAAAUA CUGAUGAG X CGAA IAUAUAU	1821
1337	AUAUCUC A UUAUAUA	686	UUAUAUA CUGAUGAG X CGAA IAGUAUA	1822
1364	AUGAUUC U CCAUAUA	687	AUAUUUG CUGAUGAG X CGAA IAAUUAU	1823
1366	GAAUCUC U AAUAUAU	688	UAAUAUU CUGAUGAG X CGAA IAGAUUC	1824
1368	AUUCUCUC U AUAUAAC	689	GUUAUAU CUGAUGAG X CGAA IAGAGAAU	1825
1370	UCUCUCUC C AUUAACUA	690	UAGUUAU CUGAUGAG X CGAA IAGAGAGA	1826
1371	CUCUCUC C UUAACUA	691	UUAGUUA CUGAUGAG X CGAA IGAGAGAG	1827
1381	AUAUAAC U AUUAGAU	692	AAUCUAAU CUGAUGAG X CGAA IUUAUAU	1828
1410	AAUGAAC U GGCCCAUC	693	GAUGGGCC CUGAUGAG X CGAA IUUCAUU	1829
1418	UUGUUGGC C UAUAACU	694	AUGUAAUA CUGAUGAG X CGAA ICCAACAA	1830
1419	UGUUGGCC C AUUACAUC	695	GAUGUAAU CUGAUGAG X CGAA IGCCAACA	1831
1420	GUUGGCC C AUUACAUC	696	AGAUGUAA CUGAUGAG X CGAA IGGCCAAC	1832
1423	GGCCCAUC U CAUCUACA	697	UGUAGAUG CUGAUGAG X CGAA IAUGGGCC	1833
1429	UCUAUAC A CAGCUGAC	698	GUCAGCUG CUGAUGAG X CGAA IUAAUAGA	1834
1432	AUAUACU U CUGACCCU	699	AGGUCAG CUGAUGAG X CGAA IAUGUAAU	1835
1435	ACAUCUAC A ACCCUUGA	700	UCAAGGU CUGAUGAG X CGAA IUAGAUGU	1836
1438	UCUACAGC U CUUGAACA	701	UGUUAAG CUGAUGAG X CGAA ICUGUAGA	1837
1442	CAGCUGAC C AACAUUGG	702	CCCAUGUU CUGAUGAG X CGAA IUCAGCUG	1838
1443	AGCUGACC C ACAUGGGG	703	CCCCAUGU CUGAUGAG X CGAA IGUCAGCU	1839
1444	GCUGACCC U CAUGGGGG	704	CCCCAUG CUGAUGAG X CGAA IGGUCAGC	1840
1450	CCUUGAAC A GGUUAGGG	705	CCCUAAC CUGAUGAG X CGAA IUUCAAGG	1841
1467	AGGGGAGC U AUUCGUGG	706	CCACGAAU CUGAUGAG X CGAA ICUCCCCU	1842
1471	GAGCUGAC A GUGGUUC	707	GGACCCAC CUGAUGAG X CGAA IUCAGCUC	1843
1483	CGUGGGUC C AAUCUUA	708	UUAAGAU CUGAUGAG X CGAA IACCCACG	1844
1486	GGGUCCGC A CUUAACUA	709	UAGUUAAG CUGAUGAG X CGAA ICGGACCC	1845
1492	GCAAAAUC U UACCUAAU	710	AUUAGGUA CUGAUGAG X CGAA IAUUUUGC	1846
1497	AUCUUAAC U AAUAGCCU	711	AGGCUAU CUGAUGAG X CGAA IUUAAGAU	1847
1500	UUAACUAC C AGCCUACU	712	AGUAGGCU CUGAUGAG X CGAA IUAGUUA	1848

Table 25

1501	UAACUACC U GCCUACUA	713	UAGUAGGC CUGAUGAG X CGAA IGUAGUUA	1849
1508	CUAAUAGC C AUUGACCA	714	UGGUCAAU CUGAUGAG X CGAA ICUAUUAG	1850
1509	UAAUAGCC U UUGACCAU	715	AUGGUCAA CUGAUGAG X CGAA IGCUAUUA	1851
1512	UAGCCUAC U ACCAUAAA	716	UUUAUGGU CUGAUGAG X CGAA IUAGGCUA	1852
1519	CUAUUGAC C ACCUACU	717	AGUAAGGU CUGAUGAG X CGAA IUCAAUAG	1853
1520	UAUUGACC A CCUACUG	718	CAGUAAGG CUGAUGAG X CGAA IGUCAAUA	1854
1526	CCAUAAAC C UGAUAACA	719	UGUUAUCA CUGAUGAG X CGAA IUUUUAGG	1855
1527	CAUAAACC U GAUAACAU	720	AUGUUAUC CUGAUGAG X CGAA IGUUUAUG	1856
1531	AACCUUAC U ACAUAAAC	721	GUUUAUGU CUGAUGAG X CGAA IUAAGGUU	1857
1538	CUGAUAAAC A CAGUAAAU	722	AUUUACUG CUGAUGAG X CGAA IUUAUCAG	1858
1544	ACAUAAAC A AUUAAACAC	723	GUGUUAUU CUGAUGAG X CGAA IUUUUAUGU	1859
1555	AAAUUAAC A UUUUGCGU	724	ACGCAAAA CUGAUGAG X CGAA IUUAAUUU	1860
1557	AUUAACAC A UUGCGUGU	725	ACACGCAA CUGAUGAG X CGAA IUGUUAUU	1861
1584	UAUUUAC A AUUCCUAC	726	GUAGGAAU CUGAUGAG X CGAA IUUAUAUA	1862
1586	UUUUAAC U UCCUACAA	727	UUGUAGGA CUGAUGAG X CGAA IUGUAUAA	1863
1593	CUAUUUUC C AUAAAGUA	728	UACUUUAU CUGAUGAG X CGAA IAAUUAUG	1864
1594	UAUAUUCC U UAAAGUAA	729	UUACUUUA CUGAUGAG X CGAA IGAAUUAU	1865
1597	AUUCCUAC A AGUAAGCU	730	AGCUUACU CUGAUGAG X CGAA IUAGGAAU	1866
1609	AAGUAAGC U AAAAUGUU	731	AACAUUUU CUGAUGAG X CGAA ICUUACUU	1867

Input Sequence = PLN. Cut Site = CH/.

Stem Length = 8 . Core Sequence = CUGAUGAG X CGAA (X = GCCGUUAGGC or other stem II)

PLN (Homo sapiens phospholamban (PLN) mRNA.; 1635 bp)

Table 26

Table 26: Human Phospholamban (PLN) G-cleaver Ribozyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
64	UCUAUACU G UGAUGAUC	732	GAUCAUCA UGAUG GCAUGCACUAUGC GCG AGUAUAGA	1868
66	UAUAUCUGU G AUGAUCAC	733	GUGAUCAU UGAUG GCAUGCACUAUGC GCG ACAGUAUA	1869
69	ACUGUGAU G AUCACAGC	734	GCUGUGAU UGAUG GCAUGCACUAUGC GCG AUCACAGU	1870
79	UCACAGCU G CCAAGGCU	735	AGCCUUGG UGAUG GCAUGCACUAUGC GCG AGCUGUGA	1871
121	AUUUGGCU G CCAGCUUU	736	AAAGCUGG UGAUG GCAUGCACUAUGC GCG AGCCAAAU	1872
143	UUUCUCUC G ACCACUUA	737	UAAGUGGU UGAUG GCAUGCACUAUGC GCG GAGAGAAA	1873
168	GACUCCCU G UCCUGCUG	738	CAGCAGGA UGAUG GCAUGCACUAUGC GCG AGGAAGUC	1874
173	CCUGUCCU G CUGGUUAC	739	GAUACCAG UGAUG GCAUGCACUAUGC GCG AGGACAGG	1875
207	CCUCACUC G CUCAGCUA	740	UAGCUGAG UGAUG GCAUGCACUAUGC GCG GAGUGAGG	1876
236	CAACCAU G AAUGCCU	741	AGGCAUUA UGAUG GCAUGCACUAUGC GCG AAUGGUUG	1877
241	AUUGAAU G CCUCAACA	742	UGUUGAGG UGAUG GCAUGCACUAUGC GCG AUUUCAAU	1878
288	CAAUUUCU G UCUCAUCU	743	AGAUGAGA UGAUG GCAUGCACUAUGC GCG AGAAUUG	1879
303	CUUAAUUAU G UCUCUUGC	744	GCAAGAGA UGAUG GCAUGCACUAUGC GCG AUUUAAG	1880
310	UGUCUCUU G CUGAUCUG	745	CAGAUACG UGAUG GCAUGCACUAUGC GCG AAGAGACA	1881
313	CUCUUGCU G AUCUGUAU	746	AUACAGAU UGAUG GCAUGCACUAUGC GCG AGCAAGAG	1882
318	GCUGAUCU G UAUCAUCG	747	CGAUGAUA UGAUG GCAUGCACUAUGC GCG AGAUCAGC	1883
328	AUCAUCGU G AUGCUUCU	748	AGAAGCAU UGAUG GCAUGCACUAUGC GCG ACGAUGAU	1884
331	AUCGUGAU G CUUCUCUG	749	CAGAGAAG UGAUG GCAUGCACUAUGC GCG AUCACGAU	1885
339	GCUCUCU G AAGUUCUG	750	CAGAACTU UGAUG GCAUGCACUAUGC GCG AGAGAAGC	1886
347	GAAGUUCU G CUACAACC	751	GGUUGUAG UGAUG GCAUGCACUAUGC GCG AGAACUUC	1887
365	CUAGAUCU G CAGCUUGC	752	GCAAGCUG UGAUG GCAUGCACUAUGC GCG AGAUCUAG	1888
372	UGCAGCUU G CCACAUA	753	UGAUGUGG UGAUG GCAUGCACUAUGC GCG AAGCUGCA	1889
392	UAAAACU G UCAUCCCA	754	UGGGAUGA UGAUG GCAUGCACUAUGC GCG AGAUUUUA	1890
402	CAUCCCAU G CAGACAGG	755	CCUGUCUG UGAUG GCAUGCACUAUGC GCG AUGGGAUG	1891
422	ACAAUAU G UAUAACAG	756	CUGUUAUA UGAUG GCAUGCACUAUGC GCG AAUAUUGU	1892
441	CACUUCU G AGUAGAAG	757	CUUCUACU UGAUG GCAUGCACUAUGC GCG AGGAAGUG	1893
459	GUUUCUUU G UGAAAAGG	758	CCUUUUA UGAUG GCAUGCACUAUGC GCG AAAGAAAC	1894
461	UUUUUUU G AAAAGGUC	759	GACUUUUU UGAUG GCAUGCACUAUGC GCG ACAAAGAA	1895

Table 26

492	AACUUAU G UUAACUA	760	UAUGGUA UGAUG GCAUGCACUAUGC GCG AAUAAGUU	1896
502	UACCAUAU G UAUAUAC	761	GAUGAAUA UGAUG GCAUGCACUAUGC GCG AU AUGGUA	1897
512	AUUAUCU G UUGGAUCU	762	AGAUCCAA UGAUG GCAUGCACUAUGC GCG AGAUGAAU	1898
522	UGGAUCUU G UAAACAUG	763	CAUGUUUA UGAUG GCAUGCACUAUGC GCG AAGAUGCA	1899
530	GUAAACAU G AAAAGGCG	764	GCCUUUU UGAUG GCAUGCACUAUGC GCG AUGUUUAC	1900
570	AAUAAGU G UAUAUAAU	765	AUUUAUA UGAUG GCAUGCACUAUGC GCG ACUAUAUU	1901
579	UAUAUAAU G CAACUGUU	766	AACAGUUG UGAUG GCAUGCACUAUGC GCG AUUUUAUA	1902
585	AUGCAACU G UUGAUUUC	767	GAUAUCAA UGAUG GCAUGCACUAUGC GCG AGUUGCAU	1903
588	CAACUGUU G AUUUCUC	768	GAGGAAAU UGAUG GCAUGCACUAUGC GCG AACAGUUG	1904
633	UCUUUUCU G AAGAUGAA	769	UUCAUCUU UGAUG GCAUGCACUAUGC GCG AGAAAGA	1905
639	CUGAAGAU G AAGAGUUU	770	AAACUCUU UGAUG GCAUGCACUAUGC GCG AUCUUCAG	1906
660	UUAAACU G CACUGCCA	771	UGGCAGUG UGAUG GCAUGCACUAUGC GCG AGUUUUAA	1907
665	ACUGCACU G CCAACAAG	772	CUUGUUGG UGAUG GCAUGCACUAUGC GCG AGUGCAGU	1908
710	ACUCUUUU G AGGUGAAU	773	AUUCACCU UGAUG GCAUGCACUAUGC GCG AAAAGAGU	1909
715	UUUGAGGU G AAUAUAAU	774	AUUAUAUU UGAUG GCAUGCACUAUGC GCG ACCUCAAA	1910
736	AUUAUAAU G UAAAGCU	775	AGCUUUUA UGAUG GCAUGCACUAUGC GCG AUUGUAAU	1911
802	ACACAAU G AAGUGUA	776	UGACACUU UGAUG GCAUGCACUAUGC GCG AUUUGUGU	1912
807	AAUGAAGU G UCAUUUAU	777	AAUAUGA UGAUG GCAUGCACUAUGC GCG ACUUCAUU	1913
830	AGUCCACU G ACUCCUCA	778	UGAGGAGU UGAUG GCAUGCACUAUGC GCG AGUGGACU	1914
844	UCACAUCU G UUAUCUUA	779	UAAGAUAU UGAUG GCAUGCACUAUGC GCG AGAUGUGA	1915
869	AACUAUUU G UAGUAACU	780	AGUUACUA UGAUG GCAUGCACUAUGC GCG AAUAAGUU	1916
907	CAGAAAUU G UAUUUUUU	781	AAAAAUUA UGAUG GCAUGCACUAUGC GCG AAUUUCUG	1917
920	UUUUCUAU G CCACAUUA	782	UAAUGUGG UGAUG GCAUGCACUAUGC GCG AUAGAAAA	1918
944	UUAAAGUU G AUGAGAAU	783	AUUCUCAU UGAUG GCAUGCACUAUGC GCG AACUUUAA	1919
947	AAGUUGAU G AGAAUCAA	784	UUUAUUUU UGAUG GCAUGCACUAUGC GCG AUCAACUU	1920
1039	UAGUACAU G UAGGUAAA	785	UUUACCUA UGAUG GCAUGCACUAUGC GCG AUGUACUA	1921
1058	AUAAUUCU G UUCUAAGA	786	UCUUAGAA UGAUG GCAUGCACUAUGC GCG AGAUUUUU	1922
1072	AGACAUUU G AUCAACAG	787	CUUUGAUU UGAUG GCAUGCACUAUGC GCG AU AUGUCU	1923
1083	CAACAGAU G AGAACUGG	788	CCAGUUUU UGAUG GCAUGCACUAUGC GCG AUCUGUUG	1924
1102	GUUAAUUAU G UGACAGUG	789	CACUGUCA UGAUG GCAUGCACUAUGC GCG AUUAUUAAC	1925
1104	UAAUAUGU G ACAGUGAG	790	CUACACUGU UGAUG GCAUGCACUAUGC GCG ACAUAUUA	1926

Table 26

1110	GUGACAGU G AGAUUAGU	791	ACUAAUCU UGAUG GCAUGCACUAUGC GCG ACUGUCAC	1927
1168	AAGGCACU G UAGUGAAU	792	AUUCACUA UGAUG GCAUGCACUAUGC GCG AGUGCCUU	1928
1173	ACUGUAGU G AAUUAUCU	793	AGAUAUUU UGAUG GCAUGCACUAUGC GCG ACUACAGU	1929
1182	AAUUAUCU G AGCUAGAG	794	CUCUAGCU UGAUG GCAUGCACUAUGC GCG AGAUAUUU	1930
1226	GGAUUCAU G AAACCUUA	795	UAAGGUUU UGAUG GCAUGCACUAUGC GCG AUGAUUCC	1931
1247	UUCAGAAU G AUUUUGCA	796	UGCAAAAU UGAUG GCAUGCACUAUGC GCG AUUCUGAA	1932
1253	AUGAUUUU G CAGGUUGU	797	ACAACCUG UGAUG GCAUGCACUAUGC GCG AAAUCAU	1933
1260	UGCAGGUU G UCUUCCAU	798	AUGGAAGA UGAUG GCAUGCACUAUGC GCG AACCUGCA	1934
1287	CAUCCAAU G CAGGCAAG	799	CUUGCCUG UGAUG GCAUGCACUAUGC GCG AUTUGAUG	1935
1316	UUUCCAGU G ACAGAAAA	800	UUUUCUGU UGAUG GCAUGCACUAUGC GCG ACUGGAAA	1936
1358	AAAUAAU G AAUUCUCU	801	AGAGAAUU UGAUG GCAUGCACUAUGC GCG AUUAUUUU	1937
1401	UAUAUUUU G AAAUGAAC	802	GUUCAUUU UGAUG GCAUGCACUAUGC GCG AAAUAUA	1938
1406	UUUGAAU G AACUUGUU	803	AACAAGUU UGAUG GCAUGCACUAUGC GCG AUUUCAAA	1939
1412	AUGAACUU G UUGGCCCA	804	UGGGCCAA UGAUG GCAUGCACUAUGC GCG AAGUUCAU	1940
1439	CUACAGCU G ACCCUUGA	805	UCAAGGGU UGAUG GCAUGCACUAUGC GCG AGCUGUAG	1941
1446	UGACCCUU G AACAUUGG	806	CCCAUGUU UGAUG GCAUGCACUAUGC GCG AAGGUCA	1942
1468	GGGAGCU G ACAAUUCG	807	CGAAUUGU UGAUG GCAUGCACUAUGC GCG AGCUCCCC	1943
1484	GUGGUCC G CAAAAUCU	808	AGAUUUUUG UGAUG GCAUGCACUAUGC GCG GGACCCAC	1944
1516	CUACUAUU G ACCAUAAA	809	UUUAUGGU UGAUG GCAUGCACUAUGC GCG AAUAGUAG	1945
1532	ACCUUACU G AUAACAUA	810	UAUGUUUU UGAUG GCAUGCACUAUGC GCG AGUAAAGU	1946
1564	CAUAUUUU G CGUGUUUU	811	AUAACACG UGAUG GCAUGCACUAUGC GCG AAAUAUUG	1947
1568	UUUUGCGU G UUAUAUGU	812	ACAUUAAA UGAUG GCAUGCACUAUGC GCG ACGCAAAA	1948
1575	UGUUUAUU G UAUUAUAC	813	GUUAUAUA UGAUG GCAUGCACUAUGC GCG AUUAUACA	1949
1619	GAGAAAAU G UUAUUUAG	814	CUAAAAUA UGAUG GCAUGCACUAUGC GCG AUUUUCUC	1950

Input Sequence = PLN. Cut Site = YG/M or UG/U.

Stem Length = 8. Core Sequence = UGAUG GCAUGCACUAUGC GCG

PLN (Homo sapiens phospholamban (PLN) mRNA.; 1635 bp)

Table 27

Table 27: Human Phospholamban (PLN) zinzyme Ribozyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
64	UCUAUACU G UGAUGAUC	732	GAUCAUCA GCCGAAAGCGAGUCAAGGUCU AGUAUAGA	1951
79	UCACAGCU G CCAAGGCU	735	AGCCUUGG GCCGAAAGCGAGUCAAGGUCU AGCUGUGA	1952
121	AUUUGGCU G CCAGCUUU	736	AAAGCUGG GCCGAAAGCGAGUCAAGGUCU AGCCAAAU	1953
168	GACUUCU G UCCUGCUG	738	CAGCAGGA GCCGAAAGCGAGUCAAGGUCU AGGAAGUC	1954
173	CCUGUCCU G CUGGUAUC	739	GAUACCAG GCCGAAAGCGAGUCAAGGUCU AGGACAGG	1955
207	CCUCACUC G CUCAGCUA	740	UAGCUGAG GCCGAAAGCGAGUCAAGGUCU GAGUGAGG	1956
241	AUUGAAAU G CCUCAACA	742	UGUUGAGG GCCGAAAGCGAGUCAAGGUCU AUUUCAAU	1957
288	CAAUUUCU G UCUCAUCU	743	AGAUGAGA GCCGAAAGCGAGUCAAGGUCU AGAAAUUG	1958
303	CUAAUAU G UCUCUUGC	744	GCAAGAGA GCCGAAAGCGAGUCAAGGUCU AUUAUUAAG	1959
310	UGUCUCU G CUGAUCUG	745	CAGAUCAG GCCGAAAGCGAGUCAAGGUCU AAGAGACA	1960
318	GCUGAUCU G UAUCAUCG	747	CGAUGAUA GCCGAAAGCGAGUCAAGGUCU AGAUCAGC	1961
331	AUCGUGAU G CUUCUCUG	749	CAGAGAAG GCCGAAAGCGAGUCAAGGUCU AUCACGAU	1962
347	GAAGUUCU G CUACAACC	751	GGUUGUAG GCCGAAAGCGAGUCAAGGUCU AGACUUC	1963
365	CUAGAUCU G CAGCUUGC	752	GCAAGCUG GCCGAAAGCGAGUCAAGGUCU AGAUCUAG	1964
372	UGCAGCUU G CCACAUCA	753	UGAUGUGG GCCGAAAGCGAGUCAAGGUCU AAGCUGCA	1965
392	UAAAACU G UCAUCCCA	754	UGGGAUGA GCCGAAAGCGAGUCAAGGUCU AGAUUUUA	1966
402	CAUCCCAU G CAGACAGG	755	CCUGUCUG GCCGAAAGCGAGUCAAGGUCU AUGGGGAG	1967
422	ACAAUAU G UAUAACAG	756	CUGUUAUA GCCGAAAGCGAGUCAAGGUCU AAUAUUUG	1968
459	GUUUCUUA G UGAAAAGG	758	CCUUUUAU GCCGAAAGCGAGUCAAGGUCU AAAGAAAC	1969
492	AACUUAU G UUAACCAUA	760	UAUGGUAA GCCGAAAGCGAGUCAAGGUCU AAUAAGUU	1970
502	UACCAUAU G UAUCAUC	761	GAUGAAUA GCCGAAAGCGAGUCAAGGUCU AUAUGGUA	1971
512	AUUGAUCU G UUGGAUCU	762	AGAUCCAA GCCGAAAGCGAGUCAAGGUCU AGAUGAAU	1972
522	UGGAUCUU G UAAACAUG	763	CAUGUUUA GCCGAAAGCGAGUCAAGGUCU AAGAUCCA	1973
570	AAUAAGU G UAUAUAUU	765	AUUUUUAU GCCGAAAGCGAGUCAAGGUCU ACUUUUUU	1974
579	UAUAAAAU G CAACUGUU	766	AACAGUUG GCCGAAAGCGAGUCAAGGUCU AUUUUAUA	1975
585	AUGCAACU G UUGAUUUC	767	GAAAUCAA GCCGAAAGCGAGUCAAGGUCU AGUUGCAU	1976
660	UUAUAACU G CACUGCCA	771	UGGCAGUG GCCGAAAGCGAGUCAAGGUCU AGUUUUAA	1977
665	ACUGCACU G CCAACAAG	772	CUUGUUGG GCCGAAAGCGAGUCAAGGUCU AGUGCAGU	1978

Table 27

736	AUUAACAAU G UAAAAGCU	775	AGUUUUU GCCGAAAGCGAGUCAAGGUCU AUUGUAAU	1979
807	AAUGAAGU G UCAUUUUU	777	AAUAAUGA GCCGAAAGCGAGUCAAGGUCU ACUUCAUU	1980
844	UCACAUCU G UUAUCUUA	779	UAAGAUAU GCCGAAAGCGAGUCAAGGUCU AGAUGUGA	1981
869	AACUAUUU G UAGUAAAU	780	AGUUAUAU GCCGAAAGCGAGUCAAGGUCU AAUAUAGU	1982
907	CAGAAAUU G UAUUUUUU	781	AAAAAUUA GCCGAAAGCGAGUCAAGGUCU AAUUUCUG	1983
920	UUUUUUUU G CCACAUUA	782	UAAUGUGG GCCGAAAGCGAGUCAAGGUCU AUAGAAAA	1984
1039	UAGUACAU G UAGGUAAA	785	UUUAACCUA GCCGAAAGCGAGUCAAGGUCU AUGUACUA	1985
1058	AUAAAUCU G UUCUAAGA	786	UCUUAGAA GCCGAAAGCGAGUCAAGGUCU AGAUUUUU	1986
1102	GUUAAUAU G UGACAGUG	789	CACUGUCA GCCGAAAGCGAGUCAAGGUCU AUUAUAAU	1987
1168	AAGGCACU G UAGUGAAU	792	AUUCACUA GCCGAAAGCGAGUCAAGGUCU AGUGCCUU	1988
1253	AUGAUUUU G CAGGUUGU	797	ACAACCUU GCCGAAAGCGAGUCAAGGUCU AAAAUCAU	1989
1260	UGCAGGUU G UCUUCCAU	798	AUGGAAGA GCCGAAAGCGAGUCAAGGUCU AACCGUCA	1990
1287	CAUCCAAU G CAGGCAAG	799	CUUGCCUG GCCGAAAGCGAGUCAAGGUCU AUUGGAUG	1991
1412	AUGAACUU G UUGGCCCA	804	UGGGCCAA GCCGAAAGCGAGUCAAGGUCU AAGUUCAU	1992
1484	GUGGUCC G CAAAUCU	808	AGAUUUUG GCCGAAAGCGAGUCAAGGUCU GGACCCAC	1993
1564	CAUAUUUU G CGUGUUUU	811	AUAACACG GCCGAAAGCGAGUCAAGGUCU AAAUAUUG	1994
1568	UUUUGCGU G UUAUAUUG	812	ACUAUAUA GCCGAAAGCGAGUCAAGGUCU ACGCAAAA	1995
1575	UGUUUAUU G UAUUAUAC	813	GUUAUAUA GCCGAAAGCGAGUCAAGGUCU AUUAUAUA	1996
1619	GAGAAAUU G UUAUUUAG	814	CUAAAUAU GCCGAAAGCGAGUCAAGGUCU AUUUUCUC	1997
21	ACUCCCCA G CUAAACAC	815	GUUUUUAG GCCGAAAGCGAGUCAAGGUCU UGGGGAGU	1998
32	AAACACCC G UAAGACUU	816	AAGUCUUA GCCGAAAGCGAGUCAAGGUCU GGGUGUUU	1999
76	UGAUCACA G CUGCCAAG	817	CUUGGCAG GCCGAAAGCGAGUCAAGGUCU UGUGAUCA	2000
85	CUGCCAAG G CUACCUAA	818	UUAGGUAG GCCGAAAGCGAGUCAAGGUCU CUUGGCAG	2001
103	AGAAGACA G UUAUCUCA	819	UGAGAUUA GCCGAAAGCGAGUCAAGGUCU UGUUUUCU	2002
118	CAUAUUUG G CUGCCAGC	820	GUUGGCAG GCCGAAAGCGAGUCAAGGUCU CAAUAUUG	2003
125	GGCUGCCA G CUUUUUUU	821	AUAAAAAG GCCGAAAGCGAGUCAAGGUCU UGGCAGCC	2004
177	UCCUGCUG G UAUCAUGG	822	CCAUGAUA GCCGAAAGCGAGUCAAGGUCU CAGCAGGA	2005
191	UGGAGAAA G UCCAAUAC	823	GUUAUGGA GCCGAAAGCGAGUCAAGGUCU UUUUCUCC	2006
212	CUCGCUCA G CUUAUAAGA	824	UCUUUAUAG GCCGAAAGCGAGUCAAGGUCU UGAGCGAG	2007
224	UAGAAGA G CCUCAACC	825	GGUUGAGG GCCGAAAGCGAGUCAAGGUCU UCUUUCUA	2008
251	CUCAACAA G CACGUCAA	826	UUACGUG GCCGAAAGCGAGUCAAGGUCU UUGUUGAG	2009

Table 27

255	ACAAAGCAC G UCAAAAGC	827	GCUUUUGA GCCGAAAGGCGAGUCAAGGUCU GUGCUUGU	2010
262	CGUCAAAA G CUACAGAA	828	UUCUGUAG GCCGAAAGGCGAGUCAAGGUCU UUUUGACG	2011
326	GUUAUC G UGAUGCUU	829	AAGCAUCA GCCGAAAGGCGAGUCAAGGUCU GAUGAUAC	2012
342	UCUCUGAA G UUCUGCUA	830	UAGCAGAA GCCGAAAGGCGAGUCAAGGUCU UUCAGAGA	2013
368	GAUCUGCA G CUUGCCAC	831	GUGGCAAG GCCGAAAGGCGAGUCAAGGUCU UGCAGAUC	2014
381	CCACAUA G CUUAAAAU	832	AUUUUAAG GCCGAAAGGCGAGUCAAGGUCU UGAUGUGG	2015
443	CUUCCUGA G UAGAAGAG	833	CUCUUCUA GCCGAAAGGCGAGUCAAGGUCU UCAGGAAG	2016
451	GUAGAAGA G UUUUUUUG	834	CAAAGAAA GCCGAAAGGCGAGUCAAGGUCU UCUUCUAC	2017
467	GUGAAAAG G UCAAGAUU	835	AAUCUUGA GCCGAAAGGCGAGUCAAGGUCU CUUUUCAC	2018
537	UGAAAAGG G CUUUUUUU	836	AAAUAAAG GCCGAAAGGCGAGUCAAGGUCU CCUUUUCA	2019
568	CAAAUAAA G UGUUAAAA	837	UUUAUACA GCCGAAAGGCGAGUCAAGGUCU UUAUUUUG	2020
603	UCAACAUG G CUCACAAA	838	UUUGUGAG GCCGAAAGGCGAGUCAAGGUCU CAUGUUGA	2021
644	GAUGAAGA G UUUAGUUU	839	AAACUAAA GCCGAAAGGCGAGUCAAGGUCU UCUTUCAUC	2022
649	AGAGUUUA G UUUUAAAA	840	UUUUAAAA GCCGAAAGGCGAGUCAAGGUCU UAAACUCU	2023
673	GCCAACAA G UUCACUUC	841	GAAGUGAA GCCGAAAGGCGAGUCAAGGUCU UUGUUGGC	2024
691	UAUAUAAA G CAUUUUUU	842	AAAUAAUG GCCGAAAGGCGAGUCAAGGUCU UUUUAUUA	2025
713	CUUUUGAG G UGAUUAUA	843	UAUAUACA GCCGAAAGGCGAGUCAAGGUCU CUCAAAAG	2026
742	AUGUAAAA G CUUCUUUA	844	UAAAGAG GCCGAAAGGCGAGUCAAGGUCU UUUUACAU	2027
758	AAUACUAA G UAUUUUUC	845	GAAAAUA GCCGAAAGGCGAGUCAAGGUCU UUAUUAUU	2028
769	UUUUUCAG G UCUUCACC	846	GGUGAAGA GCCGAAAGGCGAGUCAAGGUCU CUGAAAAA	2029
780	UUCACCAA G UAUCAAAAG	847	CUUUGAUA GCCGAAAGGCGAGUCAAGGUCU UUGGUGAA	2030
788	GUUCAA G UAAUAACA	848	UGUUUAUA GCCGAAAGGCGAGUCAAGGUCU UUUUGAUAC	2031
805	CAAAUGAA G UGUCAUUA	849	UAAUGACA GCCGAAAGGCGAGUCAAGGUCU UUCAUUUG	2032
823	UCAAAUA G UCCACUGA	850	UCAGUGGA GCCGAAAGGCGAGUCAAGGUCU UAUUUUGA	2033
872	UAUUUGUA G UAACUAUC	851	GAUAGUUA GCCGAAAGGCGAGUCAAGGUCU UACAAAUUA	2034
941	CUUUUAAA G UUGAUGAG	852	CUCAUCA GCCGAAAGGCGAGUCAAGGUCU UUUAAAAAG	2035
956	AGAAUCAA G UAUGGAAA	853	UUUCCAU GCCGAAAGGCGAGUCAAGGUCU UUGAUUCU	2036
966	AUGAAAAA G UAAGGCCA	854	UGGCCUUA GCCGAAAGGCGAGUCAAGGUCU UUUUCCAU	2037
971	AAAGUAAAG G CCAUACUC	855	GAGUAUG GCCGAAAGGCGAGUCAAGGUCU CUUACUUU	2038
1003	CCUUUUAA G UAAUUUUU	856	AAAAAUUA GCCGAAAGGCGAGUCAAGGUCU UUA AAAAGG	2039
1033	GAUUUCUA G UACAUGUA	857	UACAUGUA GCCGAAAGGCGAGUCAAGGUCU UAGAAUUC	2040

Table 27

1043	ACAUGUAG G UAAUUAU	858	AUGAUUUA GCCGAAAGCGAGUCAAGGUCU CUACAUGU	2041
1091	GAGAACUG G UGGUUAU	859	AUUAACCA GCCGAAAGCGAGUCAAGGUCU CAGUUCUC	2042
1094	AACUGGUG G UUAUAUG	860	CAUAUUA GCCGAAAGCGAGUCAAGGUCU CACCAGUU	2043
1108	AUGUGACA G UGAGAUUA	861	UAAUCUCA GCCGAAAGCGAGUCAAGGUCU UGUCACAU	2044
1117	UGAGAUUA G UCAUAUCA	862	UGAUUAUG GCCGAAAGCGAGUCAAGGUCU UAAUCUCA	2045
1163	CAUUAAG G CACUGUAG	863	CUACAGUG GCCGAAAGCGAGUCAAGGUCU CUUAAAUG	2046
1171	GCACUGUA G UGAAUUAU	864	AUAAUUA GCCGAAAGCGAGUCAAGGUCU UACAGUGC	2047
1184	UUAUCUGA G CUAGAGUU	865	AACUCUAG GCCGAAAGCGAGUCAAGGUCU UCAGAUAA	2048
1190	GAGCUAGA G UUACCUAG	866	CUAGGUAA GCCGAAAGCGAGUCAAGGUCU UCUAGCUC	2049
1198	GUUACCUA G CUUACCAU	867	AUGGUUAG GCCGAAAGCGAGUCAAGGUCU UAGGUUAC	2050
1257	UUUUGCAG G UUGUCUUC	868	GAAGACAA GCCGAAAGCGAGUCAAGGUCU CUGCAAAA	2051
1273	CCAUUCCA G CCUAACAU	869	AUGUUVAG GCCGAAAGCGAGUCAAGGUCU UGGAUUGG	2052
1291	CAAUGCAG G CAAGGAAA	870	UUUCCUUG GCCGAAAGCGAGUCAAGGUCU CUGCAUUG	2053
1314	GAUJUCCA G UGACAGAA	871	UUCUGUCA GCCGAAAGCGAGUCAAGGUCU UGGAUAUC	2054
1339	UAUCUCAA G UAUUUUU	872	AAAAAUA GCCGAAAGCGAGUCAAGGUCU UUGAGAUU	2055
1416	ACUUGUUG G CCCAUUA	873	UAGAUGGG GCCGAAAGCGAGUCAAGGUCU CAACAAGU	2056
1436	CAUCUACA G CUGACCCU	874	AGGUUCAG GCCGAAAGCGAGUCAAGGUCU UGUAGAUG	2057
1456	ACAUGGGG G UUAGGGGA	875	UCCCCUAA GCCGAAAGCGAGUCAAGGUCU CCCCAUGU	2058
1465	UUAGGGGA G CUGACAAU	876	AUUGUCAG GCCGAAAGCGAGUCAAGGUCU UCCCCUAA	2059
1476	GACAAUUC G UGGGUCCG	877	CGGACCCA GCCGAAAGCGAGUCAAGGUCU GAAUUGUC	2060
1480	AUUCGUGG G UCCGCAAA	878	UUUCCGGA GCCGAAAGCGAGUCAAGGUCU CCACGAUU	2061
1506	ACCUAAUA G CCUACUUA	879	AUAGUAGG GCCGAAAGCGAGUCAAGGUCU UAUUAGGU	2062
1545	CAUAAACA G UAAAUUAA	880	UUAUUUUA GCCGAAAGCGAGUCAAGGUCU UGUUUUUG	2063
1566	UAUUUGC G UGUUAUUA	881	AUAUAACA GCCGAAAGCGAGUCAAGGUCU GCAAAUAU	2064
1603	ACAAUAAA G UAAGCUAG	882	CUAGCUUA GCCGAAAGCGAGUCAAGGUCU UUUUUGU	2065
1607	UAAAGUAA G CUAGAGAA	883	UUCUCUAG GCCGAAAGCGAGUCAAGGUCU UUAUUUUA	2066

Input Sequence = PLN. Cut Site = G/Y

Stem Length = 8 . Core Sequence = GCcgaagGCGaGuCaaGuCu

PLN (Homo sapiens phospholamban (PLN) mRNA; 1635 bp)

Table 28

Table 28: Human Phospholamban (PLN) DNzyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
44	GACUUCAU A CAACACAA	6	TTGTGTTG GGCTAGCTACAACGA ATGAAGTC	2067
54	AACACAAU A CUCUAUAC	7	GTATAGAG GGCTAGCTACAACGA ATTGTGTT	2068
59	AAUACUCU A UACUGUGA	9	TCACAGTA GGCTAGCTACAACGA AGAGTATT	2069
61	UACUCUAU A CUGUGAUG	10	CATCACAG GGCTAGCTACAACGA ATAGAGTA	2070
88	CCAAGGCU A CCUAAAAG	12	CTTTTAGG GGCTAGCTACAACGA AGCCTTGG	2071
106	AGACAGUU A UCUCUAU	15	ATATGAGA GGCTAGCTACAACGA AACTGTCT	2072
113	UAUCUCAU A UUUGGCUG	18	CAGCCAAA GGCTAGCTACAACGA ATGAGATA	2073
132	AGCUUUUU A UCUUUCUC	25	GAGAAAAG GGCTAGCTACAACGA AAAAAGCT	2074
179	CUGCUGGU A UCAUGGAG	39	CTCCATGA GGCTAGCTACAACGA ACCAGCAG	2075
198	AGUCCAAU A CCUCACUC	42	GAGTGAGG GGCTAGCTACAACGA ATTGGACT	2076
215	GCUCAGCU A UAGAAGA	46	TCTTCTTA GGCTAGCTACAACGA AGCTGAGC	2077
265	CAAAAGCU A CAGAAUCU	52	AGATTCTG GGCTAGCTACAACGA AGCTTTTG	2078
274	CAGAAUCU A UUAUCAA	54	TTGATAAA GGCTAGCTACAACGA AGATTCTG	2079
278	AUCUAUUU A UCAAUUC	57	GAAATTGA GGCTAGCTACAACGA AAATAGAT	2080
301	AUCUAAU A UGUCUCUU	67	AAGAGACA GGCTAGCTACAACGA ATTAAGAT	2081
320	UGAUCUGU A UCAUCGUG	72	CACGATGA GGCTAGCTACAACGA ACAGATCA	2082
350	GUUCUGCU A CAACCUCU	80	AGAGGTTG GGCTAGCTACAACGA AGCAGAAC	2083
419	AAAACAAU A UUGUAUAA	91	TTATACAA GGCTAGCTACAACGA ATTGTTTT	2084
424	AAUAUUGU A UAACAGAC	93	GTCTGTTA GGCTAGCTACAACGA ACAATATT	2085
489	UAAAACUU A UUGUUACC	108	GGTAACAA GGCTAGCTACAACGA AAGTTTTA	2086
495	UUAUUGUU A CCAUAUGU	111	ACATATGG GGCTAGCTACAACGA AACAATAA	2087
500	GUUACCAU A UGUUUUCA	112	TGAATACA GGCTAGCTACAACGA ATGGAAC	2088
504	CCAUAUGU A UUCAUCUG	113	CAGATGAA GGCTAGCTACAACGA ACATATGG	2089
542	AGGGCUUU A UUUUCAA	123	TTTGAAAA GGCTAGCTACAACGA AAAGCCCT	2090
572	AUAAGUGU A UAAAAUGC	133	GCATTTTA GGCTAGCTACAACGA AACTTAT	2091
617	AAAUUUCU A UCCCAAU	144	ATTGTTGA GGCTAGCTACAACGA AGAAATTT	2092
684	CACUUCAU A UAUAAAGC	162	GCTTTATA GGCTAGCTACAACGA ATGAAGTG	2093
686	CUUCAUUA A UAAAGCAU	163	ATGCTTTA GGCTAGCTACAACGA ATATGAAG	2094
696	AAAGCAUU A UUUUUACU	166	AGTAAAAA GGCTAGCTACAACGA AATGCTTT	2095
702	UUAUUUUU A CUCUUUG	171	CAAAAGAG GGCTAGCTACAACGA AAAAATAA	2096
719	AGGUGAAU A UAAUUUAU	176	ATAAATTA GGCTAGCTACAACGA ATTACCT	2097
726	UAUAUUUU A UAUAACAA	180	TTGTAATA GGCTAGCTACAACGA AAATTATA	2098
728	UAAUUUAU A UUACAAUG	181	CATTGTAA GGCTAGCTACAACGA ATAAATTA	2099
731	UUUAUAUU A CAAUGUAA	183	TTACATTG GGCTAGCTACAACGA AATATAAA	2100
753	UCUUUAAU A CUAAGUAU	190	ATACTTAG GGCTAGCTACAACGA ATTAAAGA	2101
760	UACUAAGU A UUUUUCAG	192	CTGAAAAA GGCTAGCTACAACGA ACTTAGTA	2102
782	CACCAAGU A UCAAAGUA	201	TACTTTGA GGCTAGCTACAACGA ACTTGGTG	2103
813	GUGUCAUU A UUCAAAAU	207	ATTTTGAA GGCTAGCTACAACGA AATGACAC	2104
847	CAUCUGUU A UCUUAUUA	216	TAATAAGA GGCTAGCTACAACGA AACAGATG	2105
852	GUUAUCUU A UUAUAAAG	219	CTTTATAA GGCTAGCTACAACGA AAGATAAC	2106
855	AUCUUAUU A UAAAGAAC	221	GTTCTTTA GGCTAGCTACAACGA AATAAGAT	2107
865	AAAGAACU A UUUGUAGU	223	ACTACAAA GGCTAGCTACAACGA AGTTCTTT	2108
878	UAGUAACU A UCAGAAUC	228	GATTCTGA GGCTAGCTACAACGA AGTTACTA	2109
888	CAGAAUCU A CAUUCUAA	231	TTAGAATG GGCTAGCTACAACGA AGATTCTG	2110

Table 28

909	GAAAUUGU A UUUUUUCU	236	AGAAAAAA GGCTAGCTACAACGA ACAATTTC	2111
918	UUUUUUCU A UGCCACAU	243	ATGTGGCA GGCTAGCTACAACGA AGAAAAAA	2112
958	AAUCAAGU A UGGAAG	253	CTTTTCCA GGCTAGCTACAACGA ACTTGATT	2113
976	AAGGCCAU A CUCUACA	255	TGTAAGAG GGCTAGCTACAACGA ATGGCCTT	2114
982	AUACUCU A CAUAAUAA	258	TTATTATG GGCTAGCTACAACGA AAGAGTAT	2115
1035	AUUCUAGU A CAUGUAGG	278	CCTACATG GGCTAGCTACAACGA ACTAGAAT	2116
1070	UAAGACAU A UGAUCAAC	287	GTTGATCA GGCTAGCTACAACGA ATGTCTTA	2117
1100	UGGUUAAU A UGUGACAG	291	CTGTCACA GGCTAGCTACAACGA ATTAACCA	2118
1122	UUAGUCAU A UCACUAAU	295	ATTAGTGA GGCTAGCTACAACGA ATGACTAA	2119
1131	UCACUAAU A UACUACA	298	TGTTAGTA GGCTAGCTACAACGA ATTAGTGA	2120
1133	ACUAAUAAU A CUAACAAC	299	GTTGTAGG GGCTAGCTACAACGA ATATTAGT	2121
1178	AGUGAAU A UCUGAGCU	311	AGCTCAGA GGCTAGCTACAACGA AATTCACT	2122
1193	CUAGAGUU A CCUAGCUU	315	AAGCTAGG GGCTAGCTACAACGA AACTCTAG	2123
1202	CCUAGCUU A CCAUACUA	318	TAGTATGG GGCTAGCTACAACGA AAGCTAGG	2124
1207	CUUACCAU A CUAUAUCU	319	AGATATAG GGCTAGCTACAACGA ATGGTAAG	2125
1210	ACCAUACU A UAUCUUUG	320	CAAAGATA GGCTAGCTACAACGA AGTATGGT	2126
1212	CAUACUAAU A UCUUUGGA	321	TCCAAAGA GGCTAGCTACAACGA ATAGTATG	2127
1327	AGAAAAAU A UAUUAUCU	345	AGATAATA GGCTAGCTACAACGA ATTTTCT	2128
1329	AAAAAUAAU A UUAUCUCA	346	TGAGATAA GGCTAGCTACAACGA ATATTTTT	2129
1332	AAUAUAUU A UCUCAGU	348	ACTTGAGA GGCTAGCTACAACGA AATATATT	2130
1341	UCUCAAGU A UUUUUUAA	351	TTAAAAAA GGCTAGCTACAACGA ACTTGAGA	2131
1354	UUAAAAAU A UAUGAAUU	358	AATTCATA GGCTAGCTACAACGA ATTTTTAA	2132
1356	AAAAAUAAU A UGAAUUCU	359	AGAATTCA GGCTAGCTACAACGA ATATTTTT	2133
1375	CUCCAAAU A UUAACUAA	365	TTAGTTAA GGCTAGCTACAACGA ATTTGGAG	2134
1386	AACUAAUU A UUAGAUUA	370	TAATCTAA GGCTAGCTACAACGA AATTAGTT	2135
1394	AUUAGAUU A UAUUUUGA	374	TCAAAATA GGCTAGCTACAACGA AATCTAAT	2136
1396	UAGAUUAU A UUUUGAAA	375	TTTCAAAA GGCTAGCTACAACGA ATAATCTA	2137
1424	GCCCAUCU A UUACAUCU	382	AGATGTAA GGCTAGCTACAACGA AGATGGGC	2138
1427	CAUCUAUU A CAUCUACA	384	TGTAGATG GGCTAGCTACAACGA AATAGATG	2139
1433	UUACAUCU A CAGCUGAC	386	GTCAGCTG GGCTAGCTACAACGA AGATGTAA	2140
1498	UCUUAACU A CCUAAUAG	396	CTATTAGG GGCTAGCTACAACGA AGTTAAGA	2141
1510	AAUAGCCU A CUUUGAC	399	GTCAATAG GGCTAGCTACAACGA AGGCTATT	2142
1513	AGCCUACU A UUGACCAU	400	ATGGTCAA GGCTAGCTACAACGA AGTAGGCT	2143
1529	UAAACCUU A CUGAUAA	404	GTTATCAG GGCTAGCTACAACGA AAGGTTTA	2144
1559	UAACACAU A UUUUGCGU	410	ACGCAAAA GGCTAGCTACAACGA ATGTGTTA	2145
1571	UGCGUGUU A UAUGUAUU	415	AATACATA GGCTAGCTACAACGA AACACGCA	2146
1573	CGUGUUAU A UGUUUUAU	416	ATAATACA GGCTAGCTACAACGA ATAACACG	2147
1577	UUUAUUGU A UUAUACAC	417	GTGTATAA GGCTAGCTACAACGA ACATATAA	2148
1580	UAUGUAUU A UACACUAA	419	ATAGTGTA GGCTAGCTACAACGA AATACATA	2149
1582	UGUAUUAU A CACUAUAA	420	ATATAGTG GGCTAGCTACAACGA ATAATACA	2150
1587	UAUACACU A UAUUCCUA	421	TAGGAATA GGCTAGCTACAACGA AGTGTATA	2151
1589	UACACUAAU A UCCUACA	422	TGTAGGAA GGCTAGCTACAACGA ATAGTGTA	2152
1595	AUAUCCU A CAAUAAAG	425	CTTTATTG GGCTAGCTACAACGA AGGAATAT	2153
1622	AAAUGUU A UUUAGAAA	430	TTTCTAAA GGCTAGCTACAACGA AACATTTT	2154
64	UCUAUACU G UGAUGAUC	732	GATCATCA GGCTAGCTACAACGA AGTATAGA	2155
79	UCACAGCU G CCAAGGCU	735	AGCCTTGG GGCTAGCTACAACGA AGCTGTGA	2156
121	AUUUGGCU G CCAGCUUU	736	AAAGCTGG GGCTAGCTACAACGA AGCCAAAT	2157

Table 28

168	GACUCCU G UCCUGCUG	738	CAGCAGGA GGCTAGCTACAACGA AGGAAGTC	2158
173	CCUGUCCU G CUGGUAUC	739	GATACCAG GGCTAGCTACAACGA AGGACAGG	2159
207	CCUCACUC G CUCAGCUA	740	TAGCTGAG GGCTAGCTACAACGA GAGTGAGG	2160
241	AUUGAAAU G CCUCAACA	742	TGTTGAGG GGCTAGCTACAACGA ATTTCAAT	2161
288	CAAUUUCU G UCUCAUCU	743	AGATGAGA GGCTAGCTACAACGA AGAAATTG	2162
303	CUUAAUUA G UCUCUUGC	744	GCAAGAGA GGCTAGCTACAACGA ATATTAAG	2163
310	UGUCUCUU G CUGAUCUG	745	CAGATCAG GGCTAGCTACAACGA AAGAGACA	2164
318	GCUGAUCU G UAUCAUCG	747	CGATGATA GGCTAGCTACAACGA AGATCAGC	2165
331	AUCGUGAU G CUUCUCUG	749	CAGAGAAG GGCTAGCTACAACGA ATCACGAT	2166
347	GAAGUUCU G CUACAACC	751	GGTTGTAG GGCTAGCTACAACGA AGAATTTC	2167
365	CUAGAUCU G CAGCUUGC	752	GCAAGCTG GGCTAGCTACAACGA AGATCTAG	2168
372	UGCAGCUU G CCACAUCA	753	TGATGTGG GGCTAGCTACAACGA AAGCTGCA	2169
392	UAAAAUCU G UCAUCCCA	754	TGGGATGA GGCTAGCTACAACGA AGATTTTA	2170
402	CAUCCCAU G CAGACAGG	755	CCTGTCTG GGCTAGCTACAACGA ATGGGATG	2171
422	ACAAUAUU G UAUAACAG	756	CTGTTATA GGCTAGCTACAACGA AATATTGT	2172
459	GUUUCUUU G UGAAAAGG	758	CCTTTTCA GGCTAGCTACAACGA AAAGAAAC	2173
492	AACUUAUU G UUAACUAU	760	TATGGTAA GGCTAGCTACAACGA AATAAGTT	2174
502	UACCAUUA G UAUUAUC	761	GATGAATA GGCTAGCTACAACGA ATATGGTA	2175
512	AUUCAUCU G UUGGAUCU	762	AGATCCAA GGCTAGCTACAACGA AGATGAAT	2176
522	UGGAUCUU G UAAACAUG	763	CATGTTTA GGCTAGCTACAACGA AAGATCCA	2177
570	AAUAUAGU G UAUAUAU	765	ATTTTATA GGCTAGCTACAACGA ACTTATTT	2178
579	UAUAAAAU G CAACUGUU	766	AACAGTTG GGCTAGCTACAACGA ATTTTATA	2179
585	AUGCAACU G UUGAUUUC	767	GAAATCAA GGCTAGCTACAACGA AGTTGCAT	2180
660	UUAAAACU G CACUGCCA	771	TGGCAGTG GGCTAGCTACAACGA AGTTTTAA	2181
665	ACUGCACU G CCAACAAG	772	CTTGTTGG GGCTAGCTACAACGA AGTGCAGT	2182
736	AUUACAAU G UAAAAGCU	775	AGCTTTTA GGCTAGCTACAACGA ATTGTAAT	2183
807	AAUGAAGU G UCAUUAUU	777	AATAATGA GGCTAGCTACAACGA ACTTCATT	2184
844	UCACAUCU G UUAUCUUA	779	TAAGATAA GGCTAGCTACAACGA AGATGTGA	2185
869	AACUAUUU G UAGUAACU	780	AGTTACTA GGCTAGCTACAACGA AAATAGTT	2186
907	CAGAAAUU G UAUUUUUU	781	AAAAAATA GGCTAGCTACAACGA AATTTCTG	2187
920	UUUUCUUA G CCACAUUA	782	TAATGTGG GGCTAGCTACAACGA ATAGAAAA	2188
1039	UAGUACAU G UAGGUAAA	785	TTTACCTA GGCTAGCTACAACGA ATGTACTA	2189
1058	AUAAAUCU G UUCUAAGA	786	TCTTAGAA GGCTAGCTACAACGA AGATTTAT	2190
1102	GUUAAUUA G UGACAGUG	789	CACTGTCA GGCTAGCTACAACGA ATATTAAC	2191
1168	AAGGCACU G UAGUGAAU	792	ATTCATA GGCTAGCTACAACGA AGTGCCTT	2192
1253	AUGAUUUU G CAGGUUGU	797	ACAACCTG GGCTAGCTACAACGA AAAATCAT	2193
1260	UGCAGGUU G UCUUCCAU	798	ATGGAAGA GGCTAGCTACAACGA AACCTGCA	2194
1287	CAUCCAAU G CAGGCAAG	799	CTTGCTCG GGCTAGCTACAACGA ATTGGATG	2195
1412	AUGAACUU G UUGGCCCA	804	TGGGCCAA GGCTAGCTACAACGA AAGTTCAT	2196
1484	GUGGGUCC G CAAAUCU	808	AGATTTTG GGCTAGCTACAACGA GGACCCAC	2197
1564	CAUAUUUU G CGUGUUAU	811	ATAACACG GGCTAGCTACAACGA AAAATATG	2198
1568	UUUUGCGU G UUAUAUGU	812	ACATATAA GGCTAGCTACAACGA ACGCAAAA	2199
1575	UGUUAUUA G UAUAUAC	813	GTATAATA GGCTAGCTACAACGA ATATAACA	2200
1619	GAGAAAAU G UUAUUUAG	814	CTAAATAA GGCTAGCTACAACGA ATTTTCTC	2201
21	ACUCCCCA G CUAAACAC	815	GTGTTTAG GGCTAGCTACAACGA TGGGGAGT	2202
32	AAACACCC G UAAGACUU	816	AAGTCTTA GGCTAGCTACAACGA GGGTGTTC	2203
76	UGAUCACA G CUGCCAAG	817	CTTGGCAG GGCTAGCTACAACGA TGTGATCA	2204

Table 28

85	CUGCCAAG G CUACCUAA	818	TTAGGTAG GGCTAGCTACAACGA CTTGGCAG	2205
103	AGAAGACA G UUAUCUCA	819	TGAGATAA GGCTAGCTACAACGA TGTCTTCT	2206
118	CAUAUUUG G CUGCCAGC	820	GCTGGCAG GGCTAGCTACAACGA CAAATATG	2207
125	GGCUGCCA G CUUUUUUAU	821	ATAAAAAG GGCTAGCTACAACGA TGGCAGCC	2208
177	UCCUGCUG G UAUCAUGG	822	CCATGATA GGCTAGCTACAACGA CAGCAGGA	2209
191	UGGAGAAA G UCCAAUAC	823	GTATTGGA GGCTAGCTACAACGA TTTCTCCA	2210
212	CUCGCUCA G CUUAUAGA	824	TCTTATAG GGCTAGCTACAACGA TGAGCGAG	2211
224	UAAGAAGA G CCUCAACC	825	GGTTGAGG GGCTAGCTACAACGA TCTTCTTA	2212
251	CUCAACAA G CACGUCAA	826	TTGACGTG GGCTAGCTACAACGA TTGTTGAG	2213
255	ACAAGCAC G UCAAAAAGC	827	GCTTTTGA GGCTAGCTACAACGA GTGCTTGT	2214
262	CGUCAAAA G CUACAGAA	828	TTCTGTAG GGCTAGCTACAACGA TTTTGACG	2215
326	GUAUCAUC G UGAUGCUU	829	AAGCATCA GGCTAGCTACAACGA GATGATAC	2216
342	UCUCUGAA G UUCUGCUA	830	TAGCAGAA GGCTAGCTACAACGA TTCAGAGA	2217
368	GAUCUGCA G CUUGCCAC	831	GTGGCAAG GGCTAGCTACAACGA TGCAGATC	2218
381	CCACAUCA G CUUAAAAU	832	ATTTTAAG GGCTAGCTACAACGA TGATGTGG	2219
443	CUUCCUGA G UAGAAGAG	833	CTCTTCTA GGCTAGCTACAACGA TCAGGAAG	2220
451	GUAGAAGA G UUUCUUUG	834	CAAAGAAA GGCTAGCTACAACGA TCTTCTAC	2221
467	GUGAAAAG G UCAAGAUU	835	AATCTTGA GGCTAGCTACAACGA CTTTTCAC	2222
537	UGAAAAGG G CUUUUUUU	836	AAATAAAG GGCTAGCTACAACGA CCTTTTCA	2223
568	CAAAAUAA G UGUUAAAA	837	TTTATACA GGCTAGCTACAACGA TTATTTTG	2224
603	UCAACAUG G CUCACAAA	838	TTTGTGAG GGCTAGCTACAACGA CATGTTGA	2225
644	GAUGAAGA G UUUAGUUU	839	AAACTAAA GGCTAGCTACAACGA TCTTCATC	2226
649	AGAGUUUA G UUUUAAAA	840	TTTAAAAA GGCTAGCTACAACGA TAAACTCT	2227
673	GCCAACAA G UUCACUUC	841	GAAGTGAA GGCTAGCTACAACGA TTGTGGGC	2228
691	UAUAUAAA G CAUUAUUU	842	AAATAATG GGCTAGCTACAACGA TTTATATA	2229
713	CUUUUGAG G UGAUAUUA	843	TATATTCA GGCTAGCTACAACGA CTCAAAAG	2230
742	AUGUAAAA G CUUCUUUA	844	TAAAGAAG GGCTAGCTACAACGA TTTTACAT	2231
758	AAUACUAA G UAUUUUUC	845	GAAAAATA GGCTAGCTACAACGA TTAGTATT	2232
769	UUUUUCAG G UCUUCACC	846	GGTGAAGA GGCTAGCTACAACGA CTGAAAAA	2233
780	UUCACCAA G UAUCAAAG	847	CTTTGATA GGCTAGCTACAACGA TTGGTGAA	2234
788	GUAUCAA G UAAUAACA	848	TGTTATTA GGCTAGCTACAACGA TTTGATAC	2235
805	CAAUGAA G UGUCAUUA	849	TAATGACA GGCTAGCTACAACGA TTCATTG	2236
823	UCAAAUA G UCCACUGA	850	TCAGTGGA GGCTAGCTACAACGA TATTTTGA	2237
872	UAUUUGUA G UAACUAUC	851	GATAGTTA GGCTAGCTACAACGA TACAAATA	2238
941	CUUUUAAA G UUGAUGAG	852	CTCATCAA GGCTAGCTACAACGA TTTAAAAG	2239
956	AGAAUCAA G UAUGGAAA	853	TTCCATA GGCTAGCTACAACGA TTGATTCT	2240
966	AUGGAAA G UAAGGCCA	854	TGGCCTTA GGCTAGCTACAACGA TTTTCCAT	2241
971	AAAGUAAG G CCAUACUC	855	GAGTATGG GGCTAGCTACAACGA CTTACTTT	2242
1003	CCUUUUAA G UAAUUUUU	856	AAAAATTA GGCTAGCTACAACGA TTAAAAGG	2243
1033	GAAUUCUA G UACAUGUA	857	TACATGTA GGCTAGCTACAACGA TAGAATTC	2244
1043	ACAUGUAG G UAAAUCAU	858	ATGATTTA GGCTAGCTACAACGA CTACATGT	2245
1091	GAGAACUG G UGGUUAU	859	ATTAACCA GGCTAGCTACAACGA CAGTTCTC	2246
1094	AACUGGUG G UUAUAUG	860	CATATTAA GGCTAGCTACAACGA CACCAGTT	2247
1108	AUGUGACA G UGAGAUUA	861	TAATCTCA GGCTAGCTACAACGA TGTCACAT	2248
1117	UGAGAUUA G UCAUAUCA	862	TGATATGA GGCTAGCTACAACGA TAATCTCA	2249
1163	CAUUUAAG G CACUGUAG	863	CTACAGTG GGCTAGCTACAACGA CTTAAATG	2250
1171	GCACUGUA G UGAAUUUAU	864	ATAATTCA GGCTAGCTACAACGA TACAGTGC	2251

Table 28

1184	UUAUCUGA G CUAGAGUU	865	AACTCTAG GGCTAGCTACAACGA TCAGATAA	2252
1190	GAGCUAGA G UUACCUAG	866	CTAGGTAA GGCTAGCTACAACGA TCTAGCTC	2253
1198	GUUACCUA G CUUACCAU	867	ATGGTAAG GGCTAGCTACAACGA TAGGTAAC	2254
1257	UUUUGCAG G UUGUCUUC	868	GAAGACAA GGCTAGCTACAACGA CTGCAAAA	2255
1273	CCAUUCCA G CUAACAU	869	ATGTTAGG GGCTAGCTACAACGA TGGAAATG	2256
1291	CAAUGCAG G CAAGGAAA	870	TTTCCTTG GGCTAGCTACAACGA CTGCATTG	2257
1314	GAUUUCCA G UGACAGAA	871	TTCTGTCA GGCTAGCTACAACGA TGGAAATC	2258
1339	UAUCUCAA G UAUUUUUU	872	AAAAATA GGCTAGCTACAACGA TTGAGATA	2259
1416	ACUUGUUG G CCCAUCUA	873	TAGATGGG GGCTAGCTACAACGA CAACAAGT	2260
1436	CAUCUACA G CUGACCCU	874	AGGGTCAG GGCTAGCTACAACGA TGTAGATG	2261
1456	ACAUGGGG G UUAGGGGA	875	TCCCCTAA GGCTAGCTACAACGA CCCCATGT	2262
1465	UUAGGGGA G CUGACAAU	876	ATTGTCAG GGCTAGCTACAACGA TCCCCTAA	2263
1476	GACAAUUC G UGGGUCCG	877	CGGACCCA GGCTAGCTACAACGA GAATTGTC	2264
1480	AUUCGUGG G UCCGCAAA	878	TTTGCGGA GGCTAGCTACAACGA CCACGAAT	2265
1506	ACCUAUAU G CCUACUUA	879	ATAGTAGG GGCTAGCTACAACGA TATTAGGT	2266
1545	CAUAAACA G UAAAUUAA	880	TTAATTTA GGCTAGCTACAACGA TGTATTATG	2267
1566	UAUUUUGC G UGUUAUAU	881	ATATAACA GGCTAGCTACAACGA GCAAAATA	2268
1603	ACAUAUAA G UAAGCUAG	882	CTAGCTTA GGCTAGCTACAACGA TTTATTGT	2269
1607	UAAAGUAA G CUAGAGAA	883	TTCTCTAG GGCTAGCTACAACGA TTACTTTA	2270
13	GUCAGAAA A CUCCCCAG	884	CTGGGGAG GGCTAGCTACAACGA TTTCTGAC	2271
26	CCAGCUAA A CACCCGUA	885	TACGGGTG GGCTAGCTACAACGA TTAGCTGG	2272
28	AGCUAAAC A CCCGUAAG	886	CTTACGGG GGCTAGCTACAACGA GTTAGCT	2273
37	CCCGUAAG A CUUCAUAC	887	GTATGAAG GGCTAGCTACAACGA CTTACGGG	2274
42	AAGACUUC A UACAACAC	888	GTGTTGTA GGCTAGCTACAACGA GAAGTCTT	2275
47	UUCAUACA A CACAAUAC	889	GTATTGTG GGCTAGCTACAACGA TGTATGAA	2276
49	CAUACAAC A CAAUACUC	890	GAGTATTG GGCTAGCTACAACGA GTTGTATG	2277
52	ACAACACA A UACUCUAU	891	ATAGAGTA GGCTAGCTACAACGA TGTGTTGT	2278
67	AUACUGUG A UGAUCACA	892	TGTGATCA GGCTAGCTACAACGA CACAGTAT	2279
70	CUGUGAUG A UCACAGCU	893	AGCTGTGA GGCTAGCTACAACGA CATCACAG	2280
73	UGAUGAUC A CAGCUGCC	894	GGCAGCTG GGCTAGCTACAACGA GATCATCA	2281
100	AAAAGAAG A CAGUUAUC	895	GATAACTG GGCTAGCTACAACGA CTTCTTTT	2282
111	GUUAUCUC A UAUUUGGC	896	GCCAAATA GGCTAGCTACAACGA GAGATAAC	2283
144	UUCUCUCG A CCACUUA	897	TTAAGTGG GGCTAGCTACAACGA CGAGAGAA	2284
147	UCUCGACC A CUUAAAAC	898	GTTTTAAG GGCTAGCTACAACGA GGTCGAGA	2285
154	CACUUAUA A CUUCAGAC	899	GTCTGAAG GGCTAGCTACAACGA TTTAAGTG	2286
161	AACUUCAG A CUUCCUGU	900	ACAGGAAG GGCTAGCTACAACGA CTGAAGTT	2287
182	CUGGUUUC A UGGAGAAA	901	TTTCTCCA GGCTAGCTACAACGA GATACCAG	2288
196	AAAGUCCA A UACCUCAC	902	GTGAGGTA GGCTAGCTACAACGA TGGACTTT	2289
203	AAUACCUC A CUCGCUCA	903	TGAGCGAG GGCTAGCTACAACGA GAGGTATT	2290
230	GAGCCUCA A CCAUUGAA	904	TTCAATGG GGCTAGCTACAACGA TGAGGCTC	2291
233	CCUCAACC A UUGAAAUG	905	CATTTCAA GGCTAGCTACAACGA GGTTGAGG	2292
239	CCAUUGAA A UGCCUCAA	906	TTGAGGCA GGCTAGCTACAACGA TTCAATGG	2293
247	AUGCCUCA A CAAGCACG	907	CGTGCTTG GGCTAGCTACAACGA TGAGGCAT	2294
253	CAACAAGC A CGUAAAAA	908	TTTTGACG GGCTAGCTACAACGA GCTTGTG	2295
270	GCUACAGA A UCUAUUUA	909	TAAATAGA GGCTAGCTACAACGA TCTGTAGC	2296
282	AUUUAUCA A UUCUGUC	910	GACAGAAA GGCTAGCTACAACGA TGATAAAT	2297
293	UCUGUCUC A UCUAAUA	911	TATTAAGA GGCTAGCTACAACGA GAGACAGA	2298

Table 28

299	UCAUCUUA A UAUGUCUC	912	GAGACATA GGCTAGCTACAACGA TAAGATGA	2299
314	UCUUGCUG A UCUGUAUC	913	GATACAGA GGCTAGCTACAACGA CAGCAAGA	2300
323	UCUGUAUC A UCGUGAUG	914	CATCACGA GGCTAGCTACAACGA GATACAGA	2301
329	UCAUCGUG A UGCUUCUC	915	GAGAAGCA GGCTAGCTACAACGA CACGATGA	2302
353	CUGCUACA A CCUCUAGA	916	TCTAGAGG GGCTAGCTACAACGA TGTAGCAG	2303
361	ACCUCUAG A UCUGCAGC	917	GCTGCAGA GGCTAGCTACAACGA CTAGAGGT	2304
375	AGCUUGCC A CAUCAGCU	918	AGCTGATG GGCTAGCTACAACGA GGCAAGCT	2305
377	CUUGCCAC A UCAGCUUA	919	TAAGCTGA GGCTAGCTACAACGA GTGGCAAG	2306
388	AGCUUAAA A UCUGUCAU	920	ATGACAGA GGCTAGCTACAACGA TTAAAGCT	2307
395	AAUCUGUC A UCCCAUGC	921	GCATGGGA GGCTAGCTACAACGA GACAGATT	2308
400	GUCAUCCC A UGCAGACA	922	TGTCTGCA GGCTAGCTACAACGA GGGATGAC	2309
406	CCAUGCAG A CAGGAAAA	923	TTTTCCTG GGCTAGCTACAACGA CTGCATGG	2310
414	ACAGGAAA A CAAUAUUG	924	CAATATTG GGCTAGCTACAACGA TTTCTGT	2311
417	GGAAAACA A UAUGUAU	925	ATACAATA GGCTAGCTACAACGA TGTTTTCC	2312
427	AUUGUAUA A CAGACCAC	926	GTGGTCTG GGCTAGCTACAACGA TATACAAT	2313
431	UAUAACAG A CCACUCC	927	GGAAAGTG GGCTAGCTACAACGA CTGTTATA	2314
434	AACAGACC A CUUCCUGA	928	TCAGGAAG GGCTAGCTACAACGA GGTCGTGT	2315
473	AGGUCAAG A UUAAGACU	929	AGTCTTAA GGCTAGCTACAACGA CTTGACCT	2316
479	AGAUUAAG A CUAAAACU	930	AGTTTTAG GGCTAGCTACAACGA CTTAATCT	2317
485	AGACUAAA A CUUAUUGU	931	ACAATAAG GGCTAGCTACAACGA TTTAGTCT	2318
498	UUGUUAAC A UAUGUAU	932	AATACATA GGCTAGCTACAACGA GGTAACAA	2319
508	AUGUAUUC A UCUGUUGG	933	CCAACAGA GGCTAGCTACAACGA GAATACAT	2320
517	UCUGUUGG A UCUUGUAA	934	TTACAAGA GGCTAGCTACAACGA CCAACAGA	2321
526	UCUUGUAA A CAUGAAAA	935	TTTTCATG GGCTAGCTACAACGA TTACAAGA	2322
528	UUGUAAAC A UGAAAAGG	936	CCTTTTCA GGCTAGCTACAACGA GTTTACAA	2323
552	UUUCAAAA A UUAACUUC	937	GAAGTTAA GGCTAGCTACAACGA TTTTGAAA	2324
556	AAAAAUUA A CUUCAAAA	938	TTTGAAAG GGCTAGCTACAACGA TAATTTTT	2325
564	ACUUCAAA A UAAGUGUA	939	TACACTTA GGCTAGCTACAACGA TTTGAAGT	2326
577	UGUAUAAA A UGCAACUG	940	CAGTTGCA GGCTAGCTACAACGA TTTATACA	2327
582	AAAAUGCA A CUGUUGAU	941	ATCAACAG GGCTAGCTACAACGA TGCATTTT	2328
589	AACUGUUG A UUUCCUCA	942	TGAGGAAA GGCTAGCTACAACGA CAACAGTT	2329
598	UUUCCUCA A CAUGGCUC	943	GAGCCATG GGCTAGCTACAACGA TGAGGAAA	2330
600	UCCUCAAC A UGGCUCAC	944	GTGAGCCA GGCTAGCTACAACGA GTTGAGGA	2331
607	CAUGGCUC A CAAAUUUC	945	GAAATTTG GGCTAGCTACAACGA GAGCCATG	2332
611	GCUCACAA A UUUCUAUC	946	GATAGAAA GGCTAGCTACAACGA TTGTGAGC	2333
624	UAUCCCAA A UCUUUUCU	947	AGAAAAGA GGCTAGCTACAACGA TTGGGATA	2334
637	UUCUGAAG A UGAAGAGU	948	ACTCTTCA GGCTAGCTACAACGA CTTCAGAA	2335
657	GUUUUAAA A CUGCACUG	949	CAGTGCAG GGCTAGCTACAACGA TTTAAAAC	2336
662	AAAACUGC A CUGCCAAC	950	GTTGGCAG GGCTAGCTACAACGA GCAGTTTT	2337
669	CACUGCCA A CAAGUUCA	951	TGAACTTG GGCTAGCTACAACGA TGGCAGTG	2338
677	ACAAGUUC A CUUCAUAU	952	ATATGAAG GGCTAGCTACAACGA GAACTTGT	2339
682	UUCACUUC A UAUAUAAA	953	TTTATATA GGCTAGCTACAACGA GAAGTGAA	2340
693	UAUAAAGC A UUAUUUUU	954	AAAAATAA GGCTAGCTACAACGA GCTTTATA	2341
717	UGAGGUGA A UAUAUUUU	955	AAATTATA GGCTAGCTACAACGA TCACCTCA	2342
722	UGAAUAUA A UUAUAUUU	956	AATATAAA GGCTAGCTACAACGA TATATTCA	2343
734	AUAUUAUA A UGUAAAAG	957	CTTTTACA GGCTAGCTACAACGA TGTAATAT	2344
751	CUUCUUUA A UACUAAGU	958	ACTTAGTA GGCTAGCTACAACGA TAAAGAAG	2345

Table 28

775	AGGUCUUC A CCAAGUAU	959	ATACTTGG GGCTAGCTACAACGA GAAGACCT	2346
791	UCAAAGUA A UAACACAA	960	TTGTGTTA GGCTAGCTACAACGA TACTTTGA	2347
794	AAGUAAUA A CACAAAUG	961	CATTTGTG GGCTAGCTACAACGA TATTACTT	2348
796	GUAUAUAC A CAAAUGAA	962	TTCATTTG GGCTAGCTACAACGA GTTATTAC	2349
800	UAACACAA A UGAAGUGU	963	ACACTTCA GGCTAGCTACAACGA TTGTGTTA	2350
810	GAAGUGUC A UUAUUCAA	964	TTGAATAA GGCTAGCTACAACGA GACACTTC	2351
820	UAUUCAAA A UAGUCCAC	965	GTGGACTA GGCTAGCTACAACGA TTTGAATA	2352
827	AAUAGUCC A CUGACUCC	966	GGAGTCAG GGCTAGCTACAACGA GGACTATT	2353
831	GUCCACUG A CUCCUCAC	967	GTGAGGAG GGCTAGCTACAACGA CAGTGGAC	2354
838	GACUCCUC A CAUCUGUU	968	AACAGATG GGCTAGCTACAACGA GAGGAGTC	2355
840	CUCCUCAC A UCUGUUAU	969	ATAACAGA GGCTAGCTACAACGA GTGAGGAG	2356
862	UAUAAAGA A CUAUUGU	970	ACAAATAG GGCTAGCTACAACGA TCTTTATA	2357
875	UUGUAGUA A CUAUCAGA	971	TCTGATAG GGCTAGCTACAACGA TACTACAA	2358
884	CUAUCAGA A UCUACAUU	972	AATGTAGA GGCTAGCTACAACGA TCTGATAG	2359
890	GAAUCUAC A UUCUAAAA	973	TTTTAGAA GGCTAGCTACAACGA GTAGATTC	2360
898	AUUCUAAA A CAGAAAUU	974	AATTCTCG GGCTAGCTACAACGA TTTAGAAT	2361
904	AAACAGAA A UUGUAUUU	975	AAATACAA GGCTAGCTACAACGA TTCTGTTT	2362
923	UCUAUGCC A CAUUAACA	976	TGTTAATG GGCTAGCTACAACGA GGCATAGA	2363
925	UAUGCCAC A UUAACAUC	977	GATGTAA GGCTAGCTACAACGA GTGGCATA	2364
929	CCACAUUA A CAUCUUUU	978	AAAAGATG GGCTAGCTACAACGA TAATGTGG	2365
931	ACAUAUAC A UCUUUUAA	979	TTAAAGAA GGCTAGCTACAACGA GTTAATGT	2366
945	UAAAGUUG A UGAGAAUC	980	GATTCTCA GGCTAGCTACAACGA CAACTTTA	2367
951	UGAUGAGA A UCAAGUAU	981	ATACTTGA GGCTAGCTACAACGA TCTCATCA	2368
974	GUAAGGCC A UACUCUUA	982	TAAGAGTA GGCTAGCTACAACGA GGCCTTAC	2369
984	ACUCUUAC A UAAUAAAA	983	TTTTATTA GGCTAGCTACAACGA GTAAGAGT	2370
987	CUUACAUU A UAAAAUUC	984	GAATTTTA GGCTAGCTACAACGA TATGTAAG	2371
992	AUAAUAAA A UUCCUUUU	985	AAAAGGAA GGCTAGCTACAACGA TTTATTAT	2372
1006	UUUAAGUA A UUUUUUCA	986	TGAAAAAA GGCTAGCTACAACGA TACTTAAA	2373
1019	UUCAAAGA A UCACAGAA	987	TTCTGTGA GGCTAGCTACAACGA TCTTTGAA	2374
1022	AAAGAAUC A CAGAAUUC	988	GAATTCTG GGCTAGCTACAACGA GATTCTTT	2375
1027	AUCACAGA A UUCUAGUA	989	TACTAGAA GGCTAGCTACAACGA TCTGTGAT	2376
1037	UCUAGUAC A UGUAGGUA	990	TACCTACA GGCTAGCTACAACGA GTACTAGA	2377
1047	GUAGGUAA A UCAUAAAU	991	ATTTATGA GGCTAGCTACAACGA TTACCTAC	2378
1050	GGUAAAUC A UAAAUCUG	992	CAGATTTA GGCTAGCTACAACGA GATTTACC	2379
1054	AAUCAUAA A UCUGUUCU	993	AGAACAGA GGCTAGCTACAACGA TTATGATT	2380
1066	GUUCUAAG A CAUAUGAU	994	ATCATATG GGCTAGCTACAACGA CTTAGAAC	2381
1068	UCUAAGAC A UAUGAUCA	995	TGATCATA GGCTAGCTACAACGA GTCTTAGA	2382
1073	GACAUUUG A UCAACAGA	996	TCTGTTGA GGCTAGCTACAACGA CATATGTC	2383
1077	UAUGAUCA A CAGAUGAG	997	CTCATCTG GGCTAGCTACAACGA TGATCATA	2384
1081	AUCAACAG A UGAGAACU	998	AGTTCTCA GGCTAGCTACAACGA CTGTTGAT	2385
1087	AGAUGAGA A CUGGUGGU	999	ACCACCAG GGCTAGCTACAACGA TCTCATCT	2386
1098	GGUGGUUA A UAUGUGAC	1000	GTCACATA GGCTAGCTACAACGA TAACCACC	2387
1105	AAUAUGUG A CAGUGAGA	1001	TCTCACTG GGCTAGCTACAACGA CACATATT	2388
1113	ACAGUGAG A UUAGUCAU	1002	ATGACTAA GGCTAGCTACAACGA CTCCTGT	2389
1120	GAUUAGUC A UAUCACUA	1003	TAGTGATA GGCTAGCTACAACGA GACTAATC	2390
1125	GUCAUAUC A CUAUAUA	1004	TATATTAG GGCTAGCTACAACGA GATATGAC	2391
1129	UAUCACUA A UAUACUAA	1005	TTAGTATA GGCTAGCTACAACGA TAGTGATA	2392

Table 28

1137	AUAUACUA A CAACAGAA	1006	TTCTGTG GGCTAGCTACAACGA TAGTATAT	2393
1140	UACUAACA A CAGAAUCU	1007	AGATTCTG GGCTAGCTACAACGA TGTAGTA	2394
1145	ACAACAGA A UCUAAUCU	1008	AGATTAGA GGCTAGCTACAACGA TCTGTTGT	2395
1150	AGAAUCUA A UCUUCAUU	1009	AATGAAGA GGCTAGCTACAACGA TAGATTCT	2396
1156	UAAUCUUC A UUAAGGC	1010	GCCTTAAA GGCTAGCTACAACGA GAAGATTA	2397
1165	UUAAGGC A CUGUAGUG	1011	CACTACAG GGCTAGCTACAACGA GCCTTAAA	2398
1175	UGUAGUGA A UUAUCUGA	1012	TCAGATAA GGCTAGCTACAACGA TCACTACA	2399
1205	AGCUUACC A UACUAUAU	1013	ATATAGTA GGCTAGCTACAACGA GGTAAGCT	2400
1221	UCUUUGGA A UCAUGAAA	1014	TTTCATGA GGCTAGCTACAACGA TCCAAAGA	2401
1224	UUGGAAUC A UGAAACCU	1015	AGGTTTCA GGCTAGCTACAACGA GATTCCAA	2402
1229	AUCAUGAA A CCUUAAGA	1016	TCTTAAGG GGCTAGCTACAACGA TTCATGAT	2403
1237	ACCUUAAG A CUUCAGAA	1017	TTCTGAAG GGCTAGCTACAACGA CTTAAGGT	2404
1245	ACUUCAGA A UGAUUUUG	1018	CAAAATCA GGCTAGCTACAACGA TCTGAAGT	2405
1248	UCAGAAUG A UUUUGCAG	1019	CTGCAAAA GGCTAGCTACAACGA CATTCTGA	2406
1267	UGUCUUC A UUCCAGCC	1020	GGCTGGAA GGCTAGCTACAACGA GGAAGACA	2407
1278	CCAGCCUA A CAUCCAAU	1021	ATTGGATG GGCTAGCTACAACGA TAGGCTGG	2408
1280	AGCCUAAC A UCCAAUGC	1022	GCATTGGA GGCTAGCTACAACGA GTTAGGCT	2409
1285	AACAUCCA A UGCAGGCA	1023	TGCCTGCA GGCTAGCTACAACGA TGGATGTT	2410
1300	CAAGGAAA A UAAAAGAU	1024	ATCTTTTA GGCTAGCTACAACGA TTCCTTG	2411
1307	AAUAAAAG A UUUCAGU	1025	ACTGGAAA GGCTAGCTACAACGA CTTTTATT	2412
1317	UUCAGUG A CAGAAAAA	1026	TTTTTCTG GGCTAGCTACAACGA CACTGGAA	2413
1325	ACAGAAAA A UAUUUUAU	1027	ATAATATA GGCTAGCTACAACGA TTTTCTGT	2414
1352	UUUUAAAA A UUAUGAA	1028	TTCATATA GGCTAGCTACAACGA TTTTAAAA	2415
1360	AUAUAUGA A UUCUCUCU	1029	AGAGAGAA GGCTAGCTACAACGA TCATATAT	2416
1373	CUCUCCAA A UAUUACU	1030	AGTTAATA GGCTAGCTACAACGA TTGGAGAG	2417
1379	AAAUUAUA A CUAAUUAU	1031	ATAATTAG GGCTAGCTACAACGA TAATATTT	2418
1383	AUUAACUA A UUAUUAGA	1032	TCTAATAA GGCTAGCTACAACGA TAGTTAAT	2419
1391	AUUAUUAG A UUAUUUUU	1033	AAATATAA GGCTAGCTACAACGA CTAATAAT	2420
1404	AUUUUGAA A UGAACUUG	1034	CAAGTTCA GGCTAGCTACAACGA TTCAAAAT	2421
1408	UGAAAUGA A CUUGUUGG	1035	CCAACAAG GGCTAGCTACAACGA TCATTTCA	2422
1420	GUUGGCC C A UCUAUUAC	1036	GTAATAGA GGCTAGCTACAACGA GGGCCAAC	2423
1429	UCUAUUAC A UCUACAGC	1037	GCTGTAGA GGCTAGCTACAACGA GTAATAGA	2424
1440	UACAGCUG A CCCUUGAA	1038	TTCAAGGG GGCTAGCTACAACGA CAGCTGTA	2425
1448	ACCCUUGA A CAUGGGGG	1039	CCCCCATG GGCTAGCTACAACGA TCAAGGGT	2426
1450	CCUUGAAC A UGGGGGUU	1040	AACCCCA GGCTAGCTACAACGA GTTCAAGG	2427
1469	GGGAGCUG A CAAUUCGU	1041	ACGAATTG GGCTAGCTACAACGA CAGCTCCC	2428
1472	AGCUGACA A UUCGUGGG	1042	CCCACGAA GGCTAGCTACAACGA TGTCAGCT	2429
1489	UCCGCAAA A UCUUACU	1043	AGTTAAGA GGCTAGCTACAACGA TTTGCGGA	2430
1495	AAAUUCUA A CUACCUAA	1044	TTAGGTAG GGCTAGCTACAACGA TAAGATTT	2431
1503	ACUACCUA A UAGCCUAC	1045	GTAGGCTA GGCTAGCTACAACGA TAGGTAGT	2432
1517	UACUAUUG A CCAUAAAC	1046	GTTTATGG GGCTAGCTACAACGA CAATAGTA	2433
1520	UAUUGACC A UAAACCUU	1047	AAGGTTTA GGCTAGCTACAACGA GGTAATA	2434
1524	GACCAUAA A CCUACUG	1048	CAGTAAGG GGCTAGCTACAACGA TTATGGTC	2435
1533	CCUACUG A UAACAUAA	1049	TTATGTTA GGCTAGCTACAACGA CAGTAAGG	2436
1536	UACUGAUA A CAUAAACA	1050	TGTTTATG GGCTAGCTACAACGA TATCAGTA	2437
1538	CUGAUAA C AUAACAGU	1051	ACTGTTTA GGCTAGCTACAACGA GTTATCAG	2438
1542	UAACAUAA A CAGUAAAU	1052	ATTTACTG GGCTAGCTACAACGA TTATGTTA	2439

Table 28

1549	AACAGUAA A UUAACACA	1053	TGTGTTAA GGCTAGCTACAACGA TTACTGTT	2440
1553	GUAAAUA A CACAUUU	1054	AATATGTG GGCTAGCTACAACGA TAATTTAC	2441
1555	AAAUUAA A CAUAUUU	1055	AAAATATG GGCTAGCTACAACGA GTTAATTT	2442
1557	AUUAACAC A UAUUUUGC	1056	GCAAAATA GGCTAGCTACAACGA GTGTTAAT	2443
1584	UAUUAUAC A CUUAUUC	1057	GAATATAG GGCTAGCTACAACGA GTATAATA	2444
1598	UUCCUACA A UAAAGUAA	1058	TTACTTTA GGCTAGCTACAACGA TGTAGGAA	2445
1617	UAGAGAAA A UGUUAUUU	1059	AAATAACA GGCTAGCTACAACGA TTTCTCTA	2446

Input Sequence = PLN. Cut Site = R/Y

Stem Length = 8 . Core Sequence = GGCTAGCTACAACGA

PLN (Homo sapiens phospholamban (PLN) mRNA.; 1635 bp)

Table 29

Table 29: Human Phospholamban (PLN) amberzyme Ribozyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
64	UCUAUACU G UGAUGAUC	732	GAUCAUCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGUAUAGA	2447
66	UAUAUCUGU G AUGAUCAC	733	GUGAUCAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACAGUAUA	2448
69	ACUGUGAU G AUCACAGC	734	GCUGUGAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUCACAGU	2449
79	UCACAGCU G CCAAGGCU	735	AGCUUGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGCUGUGA	2450
121	AUUUGGCU G CCAGCUUU	736	AAAGCUGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGCCAAAU	2451
143	UUUCUCUC G ACCACUUA	737	UAAGUGGU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG GAGAGAAA	2452
168	GACUUCU G UCCUGCUG	738	CAGCAGGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGGAAGUC	2453
173	CCUGUCCU G CUGGUUAC	739	GAUACCAAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGGACAGG	2454
207	CCUCACUC G CUCAGCUA	740	UAGCUGAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG GAGUGAGG	2455
236	CAACCAUU G AAAUGCCU	741	AGGCAUUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAUGGUUG	2456
241	AUUGAAU G CCUCAACA	742	UGUGAGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUTUCAAU	2457
288	CAAUUCU G UCUCUUCU	743	AGAUGAGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGAAAUUG	2458
303	CUUAUAU G UCUCUUGC	744	GCAAGAGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUAUUAAG	2459
310	UGUCUCUU G CUGAUCUG	745	CAGAUCAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAGAGACA	2460
313	CUCUUGCU G AUCUGUAU	746	AUACAGAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGCAAGAG	2461
318	GCUGAUCU G UAUCAUCG	747	CGAUGAUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGAUCAGC	2462
328	AUCAUCGU G AUGCUUCU	748	AGAAGCAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACGAUGAU	2463
331	AUCGUGAU G CUUCUCUG	749	CAGAGAAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUCACGAU	2464
339	GCUCUCU G AAGUUCUG	750	CAGAACTU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGAGAAGC	2465
347	GAAGUUCU G CUACAACC	751	GGUUGUAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGAACUUC	2466
365	CUAGAUCU G CAGCUUGC	752	GCAAGCUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGAUCUAG	2467
372	UGCAGCUU G CCACAUCA	753	UGAUGUGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAGCUGCA	2468
392	UAAAACU G UCAUCCCA	754	UGGGAUGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGAUUUUA	2469
402	CAUCCCAU G CAGACAGG	755	CCUGUCUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUGGGAUG	2470
422	ACAAUAUU G UAUAAACAG	756	CUGUUUAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAUAUUGU	2471
441	CACUUCU G AGUAGAAG	757	CUUCUACU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGGAAGUG	2472
459	GUUUCUUU G UGAAAAGG	758	CCUUUUAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAAGAAAC	2473
461	UUUUUUUU G AAAAGGUC	759	GACCUUUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACAAAGAA	2474

Table 29

492	AACUUAU G UUAACUA	760	UAUGGUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAUAAGUU	2475
502	UACCAUAU G UAUTUAUC	761	GAUGAAUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAUAAGUA	2476
512	AUUAUAU G UUGGAUCU	762	AGAUCCAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGAUGAAU	2477
522	UGGAUCUU G UAAACAUG	763	CAUGUUUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGAUAUCC	2478
530	GUAAACAU G AAAAGGGC	764	GCCCUUUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUGUUUAC	2479
570	AAUAAGU G UAUAUAUU	765	AUUUAUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACUUUAUU	2480
579	UAUAUAU G CAACUGUU	766	AAACUUUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUUUUAUA	2481
585	AUGCAACU G UUGAUUUC	767	GAUAUCAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGUUGCAU	2482
588	CAACUGUU G AUUUCCUC	768	GAGGAAAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AACAGUUG	2483
633	UCUUUUUU G AAGAUGAA	769	UUUAUUUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGAAUAAGA	2484
639	CUGAAGAU G AAGAGUUU	770	AAACUUUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUCUUUCAG	2485
660	UAUAAACU G CACUGCCA	771	UGGCAGUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGUUUUAA	2486
665	ACUGCACU G CCAACAAG	772	CUUGUUUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGUGCAGU	2487
710	ACUCUUUU G AGGUGAAU	773	AUUCACCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAAAGAGU	2488
715	UUUGAGGU G AAUAUAUU	774	AUUUAUUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACCUCAAA	2489
736	AUUACAAU G UAAAGGCU	775	AGCUUUUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUUGUAAU	2490
802	ACACAAAU G AAGUGUCA	776	UGACACUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUUUUGUG	2491
807	AAUGAAGU G UCAUUUAU	777	AAUAUAUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACUUCAUU	2492
830	AGUCCACU G ACUCCUCA	778	UGAGGAGU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGUGGACU	2493
844	UCACAUCU G UUAUCUUA	779	UAAGAUAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGAUGUGA	2494
869	AACUAUUU G UAGUAACU	780	AGUUACUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAUAUAGU	2495
907	CAGAAAUU G UAUUUUUU	781	AAAAAUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAUUUCUG	2496
920	UUUUUUUU G CCACAUIA	782	UUAUUGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUAGAAUA	2497
944	UUAAAGUU G AUGAGAAU	783	AUUCUCAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AACUUUAA	2498
947	AAGUUGAU G AGAAUCAU	784	UUGAUUUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUCAACUU	2499
1039	UAGUACAU G UAGGUAAA	785	UUUACCUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUGUACUA	2500
1058	AUAAAUUU G UUCUAAGA	786	UCUUAGAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGAUUUUU	2501
1072	AGACAUUU G AUCAACAG	787	CUGUUGAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUUGUUCU	2502
1083	CAACAGAU G AGAACUGG	788	CCAGUUUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUCUGUUG	2503
1102	GUUAUAUU G UGACAGUG	789	CACUGUCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUUAUUAAC	2504
1104	UAUAUAUU G ACAGUGAG	790	CUACAGUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACAUAUUA	2505

Table 29

1110	GUGACAGU G AGAUUAGU	791	ACUAAUCU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG ACUGUCAC	2506
1168	AAGGCACU G UAGUGAAU	792	AUUCACUA GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AGUGCCUU	2507
1173	ACUGUAGU G AAUUAUCU	793	AGAUAUUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG ACUACAGU	2508
1182	AAUUAUCU G AGCUAGAG	794	CUCUAGCU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AGAUAUUU	2509
1226	GGAAUCAU G AAACCUUA	795	UAAGGUUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AUGAUUCC	2510
1247	UUCAGAAU G AUUUUGCA	796	UGCAAAU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AUTUCUGAA	2511
1253	AUGAUUUU G CAGGUUGU	797	ACAACCUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AAAUCAU	2512
1260	UGCAGGUU G UCUUCCAU	798	AUGGAAGA GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AACUUGCA	2513
1287	CAUCCAAU G CAGGCAAG	799	CUUGCCUG GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AUUGAUG	2514
1316	UUUCCAGU G ACAGAAAA	800	UUUUCUGU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG ACUGGAAA	2515
1358	AAAUAAU G AAUUCUCU	801	AGAGAAU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AAUAUUU	2516
1401	UAUAUUUU G AAUAUAAC	802	GUUCAUUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AAAUAUA	2517
1406	UUUGAAU G AACUUUGU	803	AACAAGUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AUUUCAAA	2518
1412	AUGAACUU G UUGGCCCA	804	UGGGCCAA GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AAGUJCAU	2519
1439	CUACAGCU G ACCCUUGA	805	UCAAGGUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AGCUUUG	2520
1446	UGACCCUU G AACAUUGG	806	CCCAUUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AAGGJUCA	2521
1468	GGGAGCU G ACAAUUCG	807	CGAUUUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AGUCCCC	2522
1484	GUGGUUCC G CAAAAUCU	808	AGAUUUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG GGACCCAC	2523
1516	CUACUAUU G ACCAUAAA	809	UUUAUGGU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AAUAGUAG	2524
1532	ACCUUACU G AUAACAUA	810	UAUGUUUU GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AGUAAAGU	2525
1564	CAUAUUUU G CGUGUUUU	811	AUAACACG GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AAAUAUUG	2526
1568	UUUGGCU G UUAUAUGU	812	ACAUUAAA GGAGAAACUCC CU UCAAGGACAUUCGUCCGG ACGCAAAA	2527
1575	UGUUAUUU G UAUUAUAC	813	GUUAUAUA GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AUAUAACA	2528
1619	GAGAAAAU G UUAUUUAG	814	CUAAUAUA GGAGAAACUCC CU UCAAGGACAUUCGUCCGG AUUUUCUC	2529
21	ACUCCCA G CUAAACAC	815	GUGUUUAG GGAGAAACUCC CU UCAAGGACAUUCGUCCGG UGGGAGU	2530
32	AAACACCC G UAAGACUU	816	AAGUCUUA GGAGAAACUCC CU UCAAGGACAUUCGUCCGG GGGUGUUU	2531
76	UGAUCACA G CUGCCAAG	817	CUUGGCAG GGAGAAACUCC CU UCAAGGACAUUCGUCCGG UGUGAUCA	2532
85	CUGCCAAG G CUACCUAA	818	UUAGGUAG GGAGAAACUCC CU UCAAGGACAUUCGUCCGG CUUGGCAG	2533
103	AGAAGACA G UUAUCUCA	819	UGAGAUAA GGAGAAACUCC CU UCAAGGACAUUCGUCCGG UGUCUUUCU	2534
118	CAUAUUUG G CUGCCAGC	820	GCUGGCAG GGAGAAACUCC CU UCAAGGACAUUCGUCCGG CAAUAUUG	2535
125	GGCUGCCA G CUUUUUUU	821	AUAAAAAG GGAGAAACUCC CU UCAAGGACAUUCGUCCGG UGGCAGCC	2536

Table 29

177	UCCUGCUG G UAUCAUGG	822	CCAUGAUA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CAGCAGGA	2537
191	UGGAGAAA G UCCAAUAC	823	GUUUGGA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUUCUCCA	2538
212	CUCGCUCA G CUAUAAGA	824	UCUUAUAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UGAGCGAG	2539
224	UAAGAAGA G CCUCAACC	825	GGUUGAGG GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UCUCUUA	2540
251	CUCAACAA G CACGUCAA	826	UUGACGUG GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUGUUGAG	2541
255	ACAAGCAC G UCAAAAGC	827	GCUUUGA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG GUGCUUGU	2542
262	CGUCARAA G CUACAGAA	828	UUCUGUAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUUGACG	2543
326	GUACAUUC G UGAUGCUU	829	AAGCAUCA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG GAUGAUAC	2544
342	UCUCUGAA G UUCUGCUA	830	UAGCAGAA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUCAGAGA	2545
368	GAUCUGCA G CUUGCCAC	831	GUGGCAAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UGCAGAUUC	2546
381	CCACAUA G CUUAAAAU	832	AUUUUAAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UGAUGUGG	2547
443	CUUCCUGA G UAGAAGAG	833	CUCUUCUA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UCAGGAAG	2548
451	GUAGAAGA G UUCUUAUG	834	CAAAGAAA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UCUCUAC	2549
467	GUGAAAAA G UCAAGAUU	835	AAUCUUGA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CUUUUCAC	2550
537	UGAAAAAG G CUUUAUUU	836	AAAUAAAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CCUUUUA	2551
568	CAAAUUA G UGUUAAAA	837	UUUAUACA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUAUUUUG	2552
603	UCAACAUG G CUCACAAA	838	UUUGUGAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CAUGUUGA	2553
644	GAUGAAGA G UUUAGUUU	839	AAACUAAA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UCUCUAC	2554
649	AGAGUUUA G UUUUAAAA	840	UUUUAAAA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UAAACUCU	2555
673	GCCAACAA G UUCACUUC	841	GAAGUGAA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUGUUGGC	2556
691	UAUAUAAA G CAUUAUUU	842	AAAUAAUG GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUUAUUA	2557
713	CUUUUGAG G UGAUAUA	843	UAUAUUCA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CUCAAAA	2558
742	AUGUAAAA G CUUCUUUA	844	UAAAGAAAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUUAACAU	2559
758	AAUACUAA G UAUUUUUC	845	GAUAAUUA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUAUUAU	2560
769	UUUUUCAG G UCUCACCC	846	GGUGAAGA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CUGAAAA	2561
780	UUCACCAA G UAUCAAAG	847	CUUUGAUA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUGGUGAA	2562
788	GUUAUCAA G UAAUAACA	848	UGUUUAUA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUUGAUAC	2563
805	CAAAUGAA G UGUCAUUA	849	UAAUGACA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUCAUUUG	2564
823	UCAAAAUA G UCCACUGA	850	UCAGUGGA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UAUUUUGA	2565
872	UAUUUGUA G UAACUAUC	851	GAUAGUUA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UACAAAA	2566
941	CUUUUAAA G UUGAUGAG	852	CUCAUCAA GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUUAAAA	2567

Table 29

956	AGAAUCAA G UAUGGAAA	853	UUUCCAUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUGAUUCU	2568
966	AUGGAAAA G UAAGGCCA	854	UGGCCUUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUUUCCA	2569
971	AAAGUAG G CCAUACUC	855	GAGUUGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CUUACUUU	2570
1003	CCUUUUA G UAAUUUUU	856	AAAAUUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUAAGG	2571
1033	GAUUCUA G UACAUGUA	857	UACAUGUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UAGAAUUC	2572
1043	ACAUGUAG G UAAAUCAU	858	AUGAUTUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CUACAUGU	2573
1091	GAGAACUG G UGGUUAUU	859	AUUAAACA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAGUUUCU	2574
1094	AACUGGUG G UUAUAUUG	860	CAUAUUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CACCAGUU	2575
1108	AUGUGACA G UGAGAUUA	861	UAAUCUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGUACAUA	2576
1117	UGAGAUUA G UCAUAUCA	862	UGAAUGA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUAUUAUG	2577
1163	CAUUUAAG G CACUGUAG	863	CUACAGUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UACAGUGC	2578
1171	GCACUGUA G UGAUUUAU	864	AUAUUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UACAGUUA	2579
1184	UUUUCUGA G CUAGAGUU	865	AACUCUAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UACAGUUA	2580
1190	GAGCUAGA G UUAACCUAG	866	CUAGGUAA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UAGGUUAC	2581
1198	GUUACCUA G CUUACCAU	867	AUGGUUAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UAGGUUAC	2582
1257	UUUUGCAG G UUGUCUUC	868	GAAGACAA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CUGCAAAA	2583
1273	CCAUUCCA G CCUAACAU	869	AUGUUAAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGGAUUGG	2584
1291	CAAUGCAG G CAAGGAAA	870	UUUCCUUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CUGCAUUG	2585
1314	GAUUUCCA G UGACAGAA	871	UUCUGUCA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGGAUUAU	2586
1339	UAUCUCAA G UAUUUUUU	872	AAAAAUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUGAGAUUA	2587
1416	ACUUGUUG G CCACUUA	873	UAGAUGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAACAAGU	2588
1436	CAUCUACA G CUGACCCU	874	AGGGUCAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGUAGAUG	2589
1456	ACAUGGGG G UUAGGGGA	875	UCCCCUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCCCAGUU	2590
1465	UUAGGGGA G CUGACAAU	876	AUUGUCAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UCCCCUUA	2591
1476	GACAAUUC G UGGGUCCG	877	CGACCCA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GAUUGUC	2592
1480	AUUCGUGG G UCCGCAAA	878	UUUGCGA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCACGAUU	2593
1506	ACCUAAUA G CCUACUUA	879	AUAGUAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UAUUAGGU	2594
1545	CAUAAACA G UAAAUUUA	880	UUAAUUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGUUUUUG	2595
1566	UAUUUUGC G UGUUAUAU	881	AUAUAACA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GCAAAAUA	2596
1603	ACAAUAAA G UAAGCUAG	882	CUAGCUUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUUUUUUG	2597
1607	UAAAGUAA G CUAGAGAA	883	UUCUCUAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUACUUUA	2598

Table 29

9	CAGAGUCA G AAAACUCC	1060	GGAGUUUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGACUCUG	2599
36	ACCCGUAA G ACUUCAU	1061	UAUGAAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUACGGGU	2600
84	GCUGCCAA G GCUACCUA	1062	UAGGUAGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUGGCAGC	2601
96	ACCUAAAA G AAGACAGU	1063	ACUGUCUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUUUAGGU	2602
99	UAAAAGAA G ACAGUUUU	1064	AUAACUGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUCUUUUA	2603
117	UCAUAUUU G GCUGCCAG	1065	CUGGCAGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AAUAUGA	2604
160	AAACUUA G ACUUCUG	1066	CAGGAAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGAAGUUU	2605
176	GUCCUGCU G GUUUAUG	1067	CAUGAUAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGCAGGAC	2606
184	GGUAUCAU G GAGAAAGU	1068	ACUUUCUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUGAUACC	2607
185	GUUAUG G AGAAAGUC	1069	GACUUUCU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAUGAUAC	2608
187	AUCAUGGA G AAAGUCCA	1070	UGGACUUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UCCAUGAU	2609
219	AGCUAUA G AAGAGCCU	1071	AGGCUCUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUAUAGCU	2610
222	UAUAAGAA G AGCCUCA	1072	UUAGAGCU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUCUUUAU	2611
268	AAGCUACA G AUUCUAUU	1073	AAUAGAUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGUAGCUU	2612
360	AACCUCUA G AUCUGCAG	1074	CUGCAGAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UAGAGGUU	2613
405	CCCAUGCA G ACAGGAA	1075	UUUCCUGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGCAUGGG	2614
409	UGCAGACA G GAAACAA	1076	UUUUUUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGUCUGCA	2615
410	GCAGACAG G AAAACAAU	1077	AUUUUUUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CUGUCUGC	2616
430	GUUAACA G ACCACUUC	1078	GAAGUGGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGUUAUAC	2617
446	CCUGAGUA G AAGAGUUU	1079	AAACUCUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UACUCAGG	2618
449	GAGUAGAA G AGUUUCUU	1080	AAGAAACU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUCUACUC	2619
466	UGUGAAAA G GUCAAGAU	1081	AUCUTUGAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUUUCACA	2620
472	AAGGUCAA G AUUAAGAC	1082	GUUUUAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUGACCUU	2621
478	AAGAUUAA G ACUAAAC	1083	GUUUUAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUAUUCUU	2622
515	CAUCUGUU G GAUCUUGU	1084	ACAAGAUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AACAGAUG	2623
516	AUCUGUUG G AUCUUGUA	1085	UACAAGAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAACAGAU	2624
535	CAUGAAAA G GGUUUUAU	1086	AUAAGCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUUUCAUG	2625
536	AUGAAAA G GCUUUUAU	1087	AAUAAAGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CUUUUCAU	2626
602	CUCAACAU G GCUCACAA	1088	UUGUGAGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUGUUGAG	2627
636	UUUCUGAA G AUGAAGAG	1089	CUCUUAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUCAGAAA	2628
642	AAGAUGAA G AGUUUAGU	1090	ACUAAACU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUCAUCUU	2629

Table 29

712	UCUUUGA G GUGAAUUAU	1091	AUAUUCAC GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UCAAAAAGA	2630
768	AUUUUUA G GUCUUCAC	1092	GUGAAGAC GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGAAAAU	2631
860	AUAUAAA G AACUAUUU	1093	AAUAGUU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UUUUAUU	2632
882	AACUAUA G AAUCUACA	1094	UGUAGAUU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGAUAGUU	2633
901	CUAAAACA G AAUUGUA	1095	UACAAUUU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGUUUUAG	2634
949	GUUGAUA G AAUCAAGU	1096	ACUUGAUU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UCAUCAAC	2635
960	UCAAGUAU G GAAAAGUA	1097	UACUUUUC GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AUACUUGA	2636
961	CAAGUAUG G AAAAGUAA	1098	UUACUUUU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG CAUACUUG	2637
970	AAAAGUAA G GCCAUACU	1099	AGUAUGGC GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UUAUUUUU	2638
1017	UUUUCAAA G AAUCACAG	1100	CUGUGAUU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UUUGAAAA	2639
1025	GAAUCACA G AAUUCUAG	1101	CUAGAAUU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGUGAUUC	2640
1042	UACAUGUA G GUAAAUAU	1102	UGAUUUAC GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UACAUGUA	2641
1065	UGUUCUAA G ACAUAUGA	1103	UCAUAUGU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UUAGAACA	2642
1080	GAUCAACA G AUGAGAAC	1104	GUUCUAU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGUUGAUC	2643
1085	ACAGAUA G AACUGGUG	1105	CACAGUU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UCAUCUGU	2644
1090	UGAGAAU G GUGGUUAA	1106	UUAACAC GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AGUUCUCA	2645
1093	GAACUGGU G GUUAAUUAU	1107	AUAUUAAC GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG ACCAGUUC	2646
1112	GACAGUA G AUUAGUCA	1108	UGACUAAU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UCAUCUGC	2647
1143	UAACAACA G AAUCUAAU	1109	AUAAGAUU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGUUGUUA	2648
1162	UCAUUUA G GCACUGUA	1110	UACAGUUC GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UUAUUAUGA	2649
1188	CUGAGCUA G AGUUACCU	1111	AGGUAAU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UAGCUCAG	2650
1218	AUAUCUUU G GAAUCAUG	1112	CAUGAUUC GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG AAAGAUAU	2651
1219	UAUCUUUG G AAUCAUGA	1113	UCAUAAU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG CAAAGAUA	2652
1236	AACCUUAA G ACUUCAGA	1114	UCUGAAGU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UUAAGGUU	2653
1243	AGACUUA G AAUGAUUU	1115	AAAUCAU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGAAGUCU	2654
1256	AUUUUGCA G GUUGUCUU	1116	AAGACAAC GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGCAAAAU	2655
1290	CCAAUGCA G GCAAGGAA	1117	UUCUUGC GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGCAUUGG	2656
1295	GCAGGCAA G GAAAUAUA	1118	UUAUUUUC GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UUGCCUUG	2657
1296	CAGGCAAG G AAAUAUAA	1119	UUUAUUUU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG CUUGCCUG	2658
1306	AAAUAAA G AUUUCAG	1120	CUGGAAU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UUUUAUUU	2659
1320	CAGUGACA G AAAAUUAU	1121	AUAUUUUU GGAGGAAACUCC CU UCAAGGACAUUCGUCCGGG UGUACUCU	2660

Table 29

1390	AAUUAUUA G AUUAUAUU	1122	AAUUAUU GGAGGAACUCC CU UCAAGGACAUCGUCCGGG UAAUAAUU	2661
1415	AACUUGUU G GCCCAUCU	1123	AGAUGGC GGAGGAACUCC CU UCAAGGACAUCGUCCGGG AACAAUUU	2662
1452	UUGAACAU G GGGUUAG	1124	CUAACCC GGAGGAACUCC CU UCAAGGACAUCGUCCGGG AUGUUCAA	2663
1453	UGAACAU G GGUUAGG	1125	CCUAAACC GGAGGAACUCC CU UCAAGGACAUCGUCCGGG CAUGUUA	2664
1454	GAACAUG G GGUUAGG	1126	CCCUAAC GGAGGAACUCC CU UCAAGGACAUCGUCCGGG CCAUGUUC	2665
1455	AACAUGG G GUUAGGG	1127	CCCUNAC GGAGGAACUCC CU UCAAGGACAUCGUCCGGG CCAUGUU	2666
1460	GGGGUUA G GGGAGCUG	1128	CAGUCCC GGAGGAACUCC CU UCAAGGACAUCGUCCGGG UAAACCCC	2667
1461	GGGUUAG G GGAGCUGA	1129	UCAGUCC GGAGGAACUCC CU UCAAGGACAUCGUCCGGG CUAACCCC	2668
1462	GGGUUAG G GAGCUGAC	1130	GUCAGCU GGAGGAACUCC CU UCAAGGACAUCGUCCGGG CCUAACCC	2669
1463	GGUAGGG G AGCUGACA	1131	UGUCAGCU GGAGGAACUCC CU UCAAGGACAUCGUCCGGG CCGUAACC	2670
1478	CAUUCGU G GGUCCGCA	1132	UGCGGACC GGAGGAACUCC CU UCAAGGACAUCGUCCGGG ACGAAUUG	2671
1479	AAUUCGU G GUCCGCAA	1133	UUGCGAC GGAGGAACUCC CU UCAAGGACAUCGUCCGGG CAGGAAUU	2672
1611	GUAAGCUA G AGAAAUUG	1134	CAUUUUU GGAGGAACUCC CU UCAAGGACAUCGUCCGGG UAGCUUAC	2673
1613	AAGCUAGA G AAAUGUU	1135	AACUUUU GGAGGAACUCC CU UCAAGGACAUCGUCCGGG UCUAGCUU	2674
1627	GUUAUUUA G AAAAUCAU	1136	AUGAUUU GGAGGAACUCC CU UCAAGGACAUCGUCCGGG UAAAUAAAC	2675

Input Sequence = PLN. Cut Site = G/.

Stem Length = 8. Core Sequence = GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG

PLN (Homo sapiens phospholamban (PLN) mRNA.; 1635 bp)

Table 30

Table 30: Human Phospholamban (PLN) Antisense and Target Sequence

Pos	Target	Seq ID	Antisense	AS Seq ID
1	CAGAGUCAGAAAACUCCCCAGCUAA	2447	TTAGCTGGGGAGTTTCTGACTCTG	3051
2	AGAGUCAGAAAACUCCCCAGCUAAA	2448	TTTAGCTGGGGAGTTTCTGACTCT	3052
3	GAGUCAGAAAACUCCCCAGCUAAAC	2449	GTTTAGCTGGGGAGTTTCTGACTC	3053
4	AGUCAGAAAACUCCCCAGCUAAACA	2450	TGTTAGCTGGGGAGTTTCTGACT	3054
5	GUCAGAAAACUCCCCAGCUAAACAC	2451	GTGTTAGCTGGGGAGTTTCTGAC	3055
6	UCAGAAAACUCCCCAGCUAAACACC	2452	GGTGTTAGCTGGGGAGTTTCTGA	3056
7	CAGAAAACUCCCCAGCUAAACACCC	2453	GGGTGTTAGCTGGGGAGTTTCTG	3057
8	AGAAAACUCCCCAGCUAAACACCCG	2454	CGGGTGTTAGCTGGGGAGTTTCT	3058
9	GAAAACUCCCCAGCUAAACACCCGU	2455	ACGGGTGTTAGCTGGGGAGTTTC	3059
10	AAAACUCCCCAGCUAAACACCCGUA	2456	TACGGGTGTTAGCTGGGGAGTTT	3060
11	AAACUCCCCAGCUAAACACCCGUA	2457	TTACGGGTGTTAGCTGGGGAGTTT	3061
12	AACUCCCCAGCUAAACACCCGUAAG	2458	CTTACGGGTGTTAGCTGGGGAGTT	3062
13	ACUCCCCAGCUAAACACCCGUAAGA	2459	TCTTACGGGTGTTAGCTGGGGAGT	3063
14	CUCCCCAGCUAAACACCCGUAAGAC	2460	GTCTTACGGGTGTTAGCTGGGGAG	3064
15	UCCCCAGCUAAACACCCGUAAGACU	2461	AGTCTTACGGGTGTTAGCTGGGGA	3065
16	CCCCAGCUAAACACCCGUAAGACUU	2462	AAGTCTTACGGGTGTTAGCTGGGG	3066
17	CCCAGCUAAACACCCGUAAGACUUC	2463	GAAGTCTTACGGGTGTTAGCTGGG	3067
18	CCAGCUAAACACCCGUAAGACUUCA	2464	TGAAGTCTTACGGGTGTTAGCTGG	3068
19	CAGCUAAACACCCGUAAGACUUCAU	2465	ATGAAGTCTTACGGGTGTTAGCTG	3069
20	AGCUAAACACCCGUAAGACUUCAUA	2466	TATGAAGTCTTACGGGTGTTAGCT	3070
21	GCUAAACACCCGUAAGACUUCAUAC	2467	GTATGAAGTCTTACGGGTGTTAGC	3071
22	CUAAACACCCGUAAGACUUCAUACA	2468	TGTATGAAGTCTTACGGGTGTTAG	3072
23	UAAACACCCGUAAGACUUCAUACAA	2469	TTGTATGAAGTCTTACGGGTGTTA	3073
24	AAACACCCGUAAGACUUCAUACAAC	2470	GTTGTATGAAGTCTTACGGGTGT	3074
25	AACACCCGUAAGACUUCAUACAACA	2471	TGTTGTATGAAGTCTTACGGGTGT	3075
26	ACACCCGUAAGACUUCAUACAACAC	2472	GTGTTGTATGAAGTCTTACGGGTGT	3076
27	CACCCGUAAGACUUCAUACAACACA	2473	TGTGTTGTATGAAGTCTTACGGGTG	3077
28	ACCCGUAAGACUUCAUACAACACAA	2474	TTGTGTTGTATGAAGTCTTACGGGT	3078
29	CCCGUAAGACUUCAUACAACACAAU	2475	ATTGTGTTGTATGAAGTCTTACGGG	3079
63	UGAUGAUCACAGCUGCCAAGGCU	2476	AGCCTTGGCAGCTGTGATCATACA	3080
64	GUGAUGAUCACAGCUGCCAAGGCUA	2477	TAGCCTTGGCAGCTGTGATCATCAC	3081
65	UGAUGAUCACAGCUGCCAAGGCUAC	2478	GTAGCCTTGGCAGCTGTGATCATCA	3082
66	GAUGAUCACAGCUGCCAAGGCUACC	2479	GGTAGCCTTGGCAGCTGTGATCATC	3083
67	AUGAUCACAGCUGCCAAGGCUACCU	2480	AGGTAGCCTTGGCAGCTGTGATCAT	3084
68	UGAUCACAGCUGCCAAGGCUACCUA	2481	TAGGTAGCCTTGGCAGCTGTGATCA	3085
69	GAUCACAGCUGCCAAGGCUACCUAA	2482	TTAGGTAGCCTTGGCAGCTGTGATC	3086
70	AUCACAGCUGCCAAGGCUACCUAAA	2483	TTTAGGTAGCCTTGGCAGCTGTGAT	3087
71	UCACAGCUGCCAAGGCUACCUAAAA	2484	TTTTAGGTAGCCTTGGCAGCTGTGA	3088
72	CACAGCUGCCAAGGCUACCUAAAAG	2485	CTTTTAGGTAGCCTTGGCAGCTGTG	3089
73	ACAGCUGCCAAGGCUACCUAAAAGA	2486	TCTTTTAGGTAGCCTTGGCAGCTGT	3090
74	CAGCUGCCAAGGCUACCUAAAAGAA	2487	TTCTTTTAGGTAGCCTTGGCAGCTG	3091
75	AGCUGCCAAGGCUACCUAAAAGAAG	2488	CTTCTTTTAGGTAGCCTTGGCAGCT	3092
76	GCUGCCAAGGCUACCUAAAAGAAGA	2489	TCTTCTTTTAGGTAGCCTTGGCAGC	3093
77	CUGCCAAGGCUACCUAAAAGAAGAC	2490	GTCTTCTTTTAGGTAGCCTTGGCAG	3094

Table 30

78	UGCCAAGGCUACCUAAAAGAAGACA	2491	TGTCTTCTTTTAGGTAGCCTTGCA	3095
79	GCCAAGGCUACCUAAAAGAAGACAG	2492	CTGTCTTCTTTTAGGTAGCCTTGGC	3096
80	CCAAGGCUACCUAAAAGAAGACAGU	2493	ACTGTCTTCTTTTAGGTAGCCTTGG	3097
81	CAAGGCUACCUAAAAGAAGACAGUU	2494	AACTGTCTTCTTTTAGGTAGCCTTG	3098
98	AGACAGUUAUCUCAUAUUUGGUGC	2495	GCAGCCAAATATGAGATAACTGTCT	3099
99	GACAGUUAUCUCAUAUUUGGUGGCC	2496	GGCAGCCAAATATGAGATAACTGTC	3100
100	ACAGUUAUCUCAUAUUUGGUGCCA	2497	TGGCAGCCAAATATGAGATAACTGT	3101
101	CAGUUAUCUCAUAUUUGGUGCCAG	2498	CTGGCAGCCAAATATGAGATAACTG	3102
102	AGUUAUCUCAUAUUUGGUGCCAGC	2499	GCTGGCAGCCAAATATGAGATAACT	3103
103	GUUAUCUCAUAUUUGGUGCCAGCU	2500	AGCTGGCAGCCAAATATGAGATAAC	3104
104	UUAUCUCAUAUUUGGUGCCAGCUU	2501	AAGCTGGCAGCCAAATATGAGATAA	3105
105	UAUCUCAUAUUUGGUGCCAGCUUU	2502	AAAGCTGGCAGCCAAATATGAGATA	3106
106	AUCUCAUAUUUGGUGCCAGCUUUU	2503	AAAAGCTGGCAGCCAAATATGAGAT	3107
107	UCUCAUAUUUGGUGCCAGCUUUUU	2504	AAAAAGCTGGCAGCCAAATATGAGA	3108
108	CUCAUAUUUGGUGCCAGCUUUUUA	2505	TAAAAAGCTGGCAGCCAAATATGAG	3109
109	UCAUAUUUGGUGCCAGCUUUUUUAU	2506	ATAAAAAGCTGGCAGCCAAATATGA	3110
110	CAUAUUUGGUGCCAGCUUUUUUAUC	2507	GATAAAAAAGCTGGCAGCCAAATATG	3111
111	AUAUUUGGUGCCAGCUUUUUUAUCU	2508	AGATAAAAAAGCTGGCAGCCAAATAT	3112
112	UAUUUGGUGCCAGCUUUUUUAUCUU	2509	AAGATAAAAAAGCTGGCAGCCAAATA	3113
113	AUUUGGUGCCAGCUUUUUUAUCUUU	2510	AAAGATAAAAAAGCTGGCAGCCAAAT	3114
114	UUUGGUGCCAGCUUUUUUAUCUUUC	2511	GAAAGATAAAAAAGCTGGCAGCCAAA	3115
115	UUGGUGCCAGCUUUUUUAUCUUUCU	2512	AGAAAGATAAAAAAGCTGGCAGCCAA	3116
116	UGGUGCCAGCUUUUUUAUCUUUCUC	2513	GAGAAAGATAAAAAAGCTGGCAGCCA	3117
117	GGCUGCCAGCUUUUUUAUCUUUCUCU	2514	AGAGAAAGATAAAAAAGCTGGCAGCC	3118
118	GCUGCCAGCUUUUUUAUCUUUCUCUC	2515	GAGAGAAAGATAAAAAAGCTGGCAGC	3119
119	CUGCCAGCUUUUUUAUCUUUCUCUG	2516	CGAGAGAAAGATAAAAAAGCTGGCAG	3120
120	UGCCAGCUUUUUUAUCUUUCUCUGA	2517	TCGAGAGAAAGATAAAAAAGCTGGCA	3121
121	GCCAGCUUUUUUAUCUUUCUCUGAC	2518	GTCGAGAGAAAGATAAAAAAGCTGGC	3122
122	CCAGCUUUUUUAUCUUUCUCUGACC	2519	GGTCGAGAGAAAGATAAAAAAGCTGG	3123
123	CAGCUUUUUUAUCUUUCUCUGACCA	2520	TGGTCGAGAGAAAGATAAAAAAGCTG	3124
124	AGCUUUUUUAUCUUUCUCUGACCAC	2521	GTGGTCGAGAGAAAGATAAAAAAGCT	3125
125	GCUUUUUUAUCUUUCUCUGACCACU	2522	AGTGGTCGAGAGAAAGATAAAAAAGC	3126
126	CUUUUUUAUCUUUCUCUGACCACUU	2523	AAGTGGTCGAGAGAAAGATAAAAAAG	3127
132	AUCUUUCUCUGACCACUUAACU	2524	AGTTTTAAGTGGTCGAGAGAAAGAT	3128
133	UCUUUCUCUGACCACUUAACUUCU	2525	AAGTTTTAAGTGGTCGAGAGAAAGA	3129
134	CUUUCUCUCUGACCACUUAACUUC	2526	GAAGTTTTAAGTGGTCGAGAGAAAG	3130
135	UUUCUCUCUGACCACUUAACUUCA	2527	TGAAGTTTTAAGTGGTCGAGAGAAA	3131
136	UUCUCUCUGACCACUUAACUUCAG	2528	CTGAAGTTTTAAGTGGTCGAGAGAA	3132
137	UCUCUCUGACCACUUAACUUCAGA	2529	TCTGAAGTTTTAAGTGGTCGAGAGA	3133
138	CUCUCUGACCACUUAACUUCAGAC	2530	GTCTGAAGTTTTAAGTGGTCGAGAG	3134
139	UCUCGACCACUUAACUUCAGACU	2531	AGTCTGAAGTTTTAAGTGGTCGAGA	3135
140	CUCGACCACUUAACUUCAGACUU	2532	AAGTCTGAAGTTTTAAGTGGTCGAG	3136
141	UCGACCACUUAACUUCAGACUUC	2533	GAAGTCTGAAGTTTTAAGTGGTCGA	3137
142	CGACCACUUAACUUCAGACUUC	2534	GGAAGTCTGAAGTTTTAAGTGGTCG	3138
143	GACCACUUAACUUCAGACUUCU	2535	AGGAAGTCTGAAGTTTTAAGTGGTC	3139
144	ACCACUUAACUUCAGACUUCUG	2536	CAGGAAGTCTGAAGTTTTAAGTGGT	3140
145	CCACUUAACUUCAGACUUCUGU	2537	ACAGGAAGTCTGAAGTTTTAAGTGG	3141

Table 30

147	ACUAAAACUUCAGACUCCUGUCC	2538	GGACAGGAAGTCTGAAGTTTAAAGT	3142
148	CUUAAAACUUCAGACUCCUGUCCU	2539	AGGACAGGAAGTCTGAAGTTTAAAG	3143
149	UUAAAACUUCAGACUCCUGUCCUG	2540	CAGGACAGGAAGTCTGAAGTTTAA	3144
150	UAAAACUUCAGACUCCUGUCCUGC	2541	GCAGGACAGGAAGTCTGAAGTTTAA	3145
151	AAAACUUCAGACUCCUGUCCUGCU	2542	AGCAGGACAGGAAGTCTGAAGTTT	3146
152	AAACUUCAGACUCCUGUCCUGCUG	2543	CAGCAGGACAGGAAGTCTGAAGTTT	3147
153	AACUUCAGACUCCUGUCCUGCUGG	2544	CCAGCAGGACAGGAAGTCTGAAGTT	3148
154	ACUUCAGACUCCUGUCCUGCUGGU	2545	ACCAGCAGGACAGGAAGTCTGAAGT	3149
155	CUUCAGACUCCUGUCCUGCUGGUA	2546	TACCAGCAGGACAGGAAGTCTGAAG	3150
156	UUCAGACUCCUGUCCUGCUGGUAU	2547	ATACCAGCAGGACAGGAAGTCTGAA	3151
157	UCAGACUCCUGUCCUGCUGGUUUC	2548	GATACCAGCAGGACAGGAAGTCTGA	3152
158	CAGACUCCUGUCCUGCUGGUUACA	2549	TGATACCAGCAGGACAGGAAGTCTG	3153
159	AGACUCCUGUCCUGCUGGUUAUCAU	2550	ATGATACCAGCAGGACAGGAAGTCT	3154
160	GACUCCUGUCCUGCUGGUUAUCAUG	2551	CATGATACCAGCAGGACAGGAAGTC	3155
161	ACUCCUGUCCUGCUGGUUAUCAUGG	2552	CCATGATACCAGCAGGACAGGAAGT	3156
162	CUCCUGUCCUGCUGGUUAUCAUGGA	2553	TCCATGATACCAGCAGGACAGGAAG	3157
163	UCCUGUCCUGCUGGUUAUCAUGGAG	2554	CTCCATGATACCAGCAGGACAGGAA	3158
164	UCCUGUCCUGCUGGUUAUCAUGGAGA	2555	TCTCCATGATACCAGCAGGACAGGA	3159
165	CCUGUCCUGCUGGUUAUCAUGGAGAA	2556	TTCTCCATGATACCAGCAGGACAGG	3160
166	CUGUCCUGCUGGUUAUCAUGGAGAAA	2557	TTTCTCCATGATACCAGCAGGACAG	3161
167	UGUCCUGCUGGUUAUCAUGGAGAAAG	2558	CTTCTCCATGATACCAGCAGGACA	3162
168	GUCCUGCUGGUUAUCAUGGAGAAAGU	2559	ACTTCTCCATGATACCAGCAGGAC	3163
169	UCCUGCUGGUUAUCAUGGAGAAAGUC	2560	GACTTCTCCATGATACCAGCAGGA	3164
170	CCUGCUGGUUAUCAUGGAGAAAGUCC	2561	GGACTTCTCCATGATACCAGCAGG	3165
180	UCAUGGAGAAAGUCCAAUACCUCAC	2562	GTGAGGTATTGGACTTTCTCCATGA	3166
181	CAUGGAGAAAGUCCAAUACCUCACU	2563	AGTGAGGTATTGGACTTTCTCCATG	3167
182	AUGGAGAAAGUCCAAUACCUCACUC	2564	GAGTGAGGTATTGGACTTTCTCCAT	3168
183	UGGAGAAAGUCCAAUACCUCACUCG	2565	CGAGTGAGGTATTGGACTTTCTCCA	3169
184	GGAGAAAGUCCAAUACCUCACUCGC	2566	GCGAGTGAGGTATTGGACTTTCTCC	3170
185	GAGAAAGUCCAAUACCUCACUCGCU	2567	AGCGAGTGAGGTATTGGACTTTCTC	3171
186	AGAAAGUCCAAUACCUCACUCGCUC	2568	GAGCGAGTGAGGTATTGGACTTTCT	3172
187	GAAAGUCCAAUACCUCACUCGCUCA	2569	TGAGCGAGTGAGGTATTGGACTTTC	3173
188	AAAGUCCAAUACCUCACUCGCUCAG	2570	CTGAGCGAGTGAGGTATTGGACTTT	3174
189	AAGUCCAAUACCUCACUCGCUCAGC	2571	GCTGAGCGAGTGAGGTATTGGACTT	3175
190	AGUCCAAUACCUCACUCGCUCAGCU	2572	AGCTGAGCGAGTGAGGTATTGGACT	3176
191	GUCCAAUACCUCACUCGCUCAGCUA	2573	TAGCTGAGCGAGTGAGGTATTGGAC	3177
192	UCCAAUACCUCACUCGCUCAGCUAU	2574	ATAGCTGAGCGAGTGAGGTATTGGA	3178
193	CCAAUACCUCACUCGCUCAGCUAUA	2575	TATAGCTGAGCGAGTGAGGTATTGG	3179
194	CAAUACCUCACUCGCUCAGCUAUAA	2576	TTATAGCTGAGCGAGTGAGGTATTG	3180
195	AAUACCUCACUCGCUCAGCUAUAAAG	2577	CTTATAGCTGAGCGAGTGAGGTATT	3181
196	AUACCUCACUCGCUCAGCUAUAAAGA	2578	TCTTATAGCTGAGCGAGTGAGGTAT	3182
197	UACCUCACUCGCUCAGCUAUAAAGAA	2579	TTCTTATAGCTGAGCGAGTGAGGTA	3183
198	ACCUCACUCGCUCAGCUAUAAAGAAG	2580	CTTCTTATAGCTGAGCGAGTGAGGT	3184
199	CCUCACUCGCUCAGCUAUAAAGAAGA	2581	TCTTCTTATAGCTGAGCGAGTGAGG	3185
200	CUCACUCGCUCAGCUAUAAAGAAGAG	2582	CTCTTCTTATAGCTGAGCGAGTGAG	3186
201	UCACUCGCUCAGCUAUAAAGAAGAGC	2583	GCTCTTCTTATAGCTGAGCGAGTGGA	3187
202	CACUCGCUCAGCUAUAAAGAAGAGCC	2584	GGCTCTTCTTATAGCTGAGCGAGTG	3188

Table 30

203	ACUCGCUAGCUAUAAGAAGAGCCU	2585	AGGCTCTTCTTATAGCTGAGCGAGT	3189
204	CUCGCUAGCUAUAAGAAGAGCCUC	2586	GAGGCTCTTCTTATAGCTGAGCGAG	3190
205	UCGCUAGCUAUAAGAAGAGCCUCA	2587	TGAGGCTCTTCTTATAGCTGAGCGA	3191
206	CGCUCAGCUAUAAGAAGAGCCUCA	2588	TTGAGGCTCTTCTTATAGCTGAGCG	3192
207	GCUCAGCUAUAAGAAGAGCCUCAAC	2589	GTTGAGGCTCTTCTTATAGCTGAGC	3193
208	CUCAGCUAUAAGAAGAGCCUCAACC	2590	GGTTGAGGCTCTTCTTATAGCTGAG	3194
209	UCAGCUAUAAGAAGAGCCUCAACCA	2591	TGTTGAGGCTCTTCTTATAGCTGA	3195
210	CAGCUAUAAGAAGAGCCUCAACCAU	2592	ATGGTTGAGGCTCTTCTTATAGCTG	3196
211	AGCUAUAAGAAGAGCCUCAACCAU	2593	AATGGTTGAGGCTCTTCTTATAGCT	3197
212	GCUAUAAGAAGAGCCUCAACCAUUG	2594	CAATGGTTGAGGCTCTTCTTATAGC	3198
213	CUAUAAGAAGAGCCUCAACCAUUGA	2595	TCAATGGTTGAGGCTCTTCTTATAG	3199
214	UAUAAGAAGAGCCUCAACCAUUGAA	2596	TTCAATGGTTGAGGCTCTTCTTATA	3200
215	AUAAGAAGAGCCUCAACCAUUGAAA	2597	TTTCAATGGTTGAGGCTCTTCTTAT	3201
216	UAAGAAGAGCCUCAACCAUUGAAAU	2598	ATTTCATGGTTGAGGCTCTTCTTA	3202
217	AAGAAGAGCCUCAACCAUUGAAAUG	2599	CATTTCAATGGTTGAGGCTCTTCTT	3203
218	AGAAGAGCCUCAACCAUUGAAAUGC	2600	GCATTTCAATGGTTGAGGCTCTTCT	3204
219	GAAGAGCCUCAACCAUUGAAAUGCC	2601	GGCATTTCAATGGTTGAGGCTCTTC	3205
220	AAGAGCCUCAACCAUUGAAAUGCCU	2602	AGGCATTTCAATGGTTGAGGCTCTT	3206
221	AGAGCCUCAACCAUUGAAAUGCCUC	2603	GAGGCATTTCAATGGTTGAGGCTCT	3207
222	GAGCCUCAACCAUUGAAAUGCCUCA	2604	TGAGGCATTTCAATGGTTGAGGCTC	3208
223	AGCCUCAACCAUUGAAAUGCCUCA	2605	TTGAGGCATTTCAATGGTTGAGGCT	3209
224	GCCUCAACCAUUGAAAUGCCUCAAC	2606	GTTGAGGCATTTCAATGGTTGAGGC	3210
225	CCUCAACCAUUGAAAUGCCUCAACA	2607	TGTTGAGGCATTTCAATGGTTGAGG	3211
226	CUCAACCAUUGAAAUGCCUCAACAA	2608	TTGTTGAGGCATTTCAATGGTTGAG	3212
227	UCAACCAUUGAAAUGCCUCAACAAG	2609	CTTGTTGAGGCATTTCAATGGTTGA	3213
228	CAACCAUUGAAAUGCCUCAACAAGC	2610	GCTTGTTGAGGCATTTCAATGGTTG	3214
229	AACCAUUGAAAUGCCUCAACAAGCA	2611	TGCTTGTTGAGGCATTTCAATGGTT	3215
230	ACCAUUGAAAUGCCUCAACAAGCAC	2612	GTGCTTGTTGAGGCATTTCAATGGT	3216
231	CCAUUGAAAUGCCUCAACAAGCACG	2613	CGTGCTTGTTGAGGCATTTCAATGG	3217
232	CAUUGAAAUGCCUCAACAAGCACGU	2614	ACGTGCTTGTTGAGGCATTTCAATG	3218
233	AUUGAAAUGCCUCAACAAGCACGUC	2615	GACGTGCTTGTTGAGGCATTTCAAT	3219
234	UUGAAAUGCCUCAACAAGCACGUCA	2616	TGACGTGCTTGTTGAGGCATTTCAA	3220
235	UGAAAUGCCUCAACAAGCACGUCAA	2617	TTGACGTGCTTGTTGAGGCATTTCA	3221
236	GAAAUGCCUCAACAAGCACGUCAAA	2618	TTTGACGTGCTTGTTGAGGCATTTT	3222
237	AAAUGCCUCAACAAGCACGUCAAAA	2619	TTTTGACGTGCTTGTTGAGGCATTT	3223
238	AAUGCCUCAACAAGCACGUCAAAAAG	2620	CTTTTGACGTGCTTGTTGAGGCATT	3224
239	AUGCCUCAACAAGCACGUCAAAAAGC	2621	GCTTTTGACGTGCTTGTTGAGGCAT	3225
240	UGCCUCAACAAGCACGUCAAAAAGCU	2622	AGCTTTTGACGTGCTTGTTGAGGCA	3226
241	GCCUCAACAAGCACGUCAAAAAGCUA	2623	TAGCTTTTGACGTGCTTGTTGAGGC	3227
242	CCUCAACAAGCACGUCAAAAAGCUAC	2624	GTAGCTTTTGACGTGCTTGTTGAGG	3228
243	CUCAACAAGCACGUCAAAAAGCUACA	2625	TGTAGCTTTTGACGTGCTTGTTGAG	3229
244	UCAACAAGCACGUCAAAAAGCUACAG	2626	CTGTAGCTTTTGACGTGCTTGTTGA	3230
245	CAACAAGCACGUCAAAAAGCUACAGA	2627	TCTGTAGCTTTTGACGTGCTTGTTG	3231
246	AACAAGCACGUCAAAAAGCUACAGAA	2628	TTCTGTAGCTTTTGACGTGCTTGTT	3232
247	ACAAGCACGUCAAAAAGCUACAGAAU	2629	ATTCTGTAGCTTTTGACGTGCTTGTT	3233
248	CAAGCACGUCAAAAAGCUACAGAAUC	2630	GATTCTGTAGCTTTTGACGTGCTTG	3234
249	AAGCACGUCAAAAAGCUACAGAAUCU	2631	AGATTCTGTAGCTTTTGACGTGCTT	3235

Table 30

250	AGCACGUCAAAGCUACAGAAUCUA	2632	TAGATTCTGTAGCTTTTGACGTGCT	3236
251	GCACGUCAAAGCUACAGAAUCUAU	2633	ATAGATTCTGTAGCTTTTGACGTGC	3237
252	CACGUCAAAGCUACAGAAUCUAUU	2634	AATAGATTCTGTAGCTTTTGACGTG	3238
253	ACGUCAAAGCUACAGAAUCUAUUU	2635	AAATAGATTCTGTAGCTTTTGACGT	3239
254	CGUCAAAGCUACAGAAUCUAUUUA	2636	TAAATAGATTCTGTAGCTTTTGACG	3240
255	GUCAAAAGCUACAGAAUCUAUUUAU	2637	ATAAATAGATTCTGTAGCTTTTGAC	3241
256	UCAAAAGCUACAGAAUCUAUUUAUC	2638	GATAAATAGATTCTGTAGCTTTTGA	3242
257	CAAAAGCUACAGAAUCUAUUUAUCA	2639	TGATAAATAGATTCTGTAGCTTTTG	3243
258	AAAAGCUACAGAAUCUAUUUAUCAA	2640	TTGATAAATAGATTCTGTAGCTTTT	3244
259	AAAGCUACAGAAUCUAUUUAUCAU	2641	ATTGATAAATAGATTCTGTAGCTTT	3245
260	AAGCUACAGAAUCUAUUUAUCAUU	2642	AATTGATAAATAGATTCTGTAGCTT	3246
261	AGCUACAGAAUCUAUUUAUCAUUU	2643	AAATTGATAAATAGATTCTGTAGCT	3247
262	GCUACAGAAUCUAUUUAUCAUUUC	2644	GAAATTGATAAATAGATTCTGTAGC	3248
263	CUACAGAAUCUAUUUAUCAUUUCU	2645	AGAAATTGATAAATAGATTCTGTAG	3249
264	UACAGAAUCUAUUUAUCAUUUCUG	2646	CAGAAATTGATAAATAGATTCTGTA	3250
265	ACAGAAUCUAUUUAUCAUUUCUGU	2647	ACAGAAATTGATAAATAGATTCTGT	3251
266	CAGAAUCUAUUUAUCAUUUCUGUC	2648	GACAGAAATTGATAAATAGATTCTG	3252
267	AGAAUCUAUUUAUCAUUUCUGUCU	2649	AGACAGAAATTGATAAATAGATTCT	3253
268	GAAUCUAUUUAUCAUUUCUGUCUC	2650	GAGACAGAAATTGATAAATAGATTCT	3254
269	AAUCUAUUUAUCAUUUCUGUCUCA	2651	TGAGACAGAAATTGATAAATAGATT	3255
270	AUCUAUUUAUCAUUUCUGUCUCAU	2652	ATGAGACAGAAATTGATAAATAGAT	3256
271	UCUAUUUAUCAUUUCUGUCUCAUC	2653	GATGAGACAGAAATTGATAAATAGA	3257
272	CUAUUUUAUCAUUUCUGUCUCAUCU	2654	AGATGAGACAGAAATTGATAAATAG	3258
273	UAUUUAUCAUUUCUGUCUCAUCUU	2655	AAGATGAGACAGAAATTGATAAATA	3259
274	AUUUAUCAUUUCUGUCUCAUCUUA	2656	TAAGATGAGACAGAAATTGATAAAT	3260
275	UUUAUCAUUUCUGUCUCAUCUUAA	2657	TTAAGATGAGACAGAAATTGATAAA	3261
276	UUAUCAUUUCUGUCUCAUCUUAAU	2658	ATTAAGATGAGACAGAAATTGATAA	3262
277	UAUCAUUUCUGUCUCAUCUUAAUA	2659	TATTAAGATGAGACAGAAATTGATA	3263
278	AUCAUUUCUGUCUCAUCUUAAUAU	2660	ATATTAAGATGAGACAGAAATTGAT	3264
279	UCAUUUCUGUCUCAUCUUAAUAUG	2661	CATATTAAGATGAGACAGAAATTGA	3265
280	CAUUUCUGUCUCAUCUUAAUAUGU	2662	ACATATTAAGATGAGACAGAAATTG	3266
281	AAUUUCUGUCUCAUCUUAAUAUGUC	2663	GACATATTAAGATGAGACAGAAATT	3267
282	AUUUCUGUCUCAUCUUAAUAUGUCU	2664	AGACATATTAAGATGAGACAGAAAT	3268
283	UUUCUGUCUCAUCUUAAUAUGUCUC	2665	GAGACATATTAAGATGAGACAGAAA	3269
284	UUCUGUCUCAUCUUAAUAUGUCUCU	2666	AGAGACATATTAAGATGAGACAGAA	3270
285	UCUGUCUCAUCUUAAUAUGUCUCUU	2667	AAGAGACATATTAAGATGAGACAGA	3271
286	CUGUCUCAUCUUAAUAUGUCUCUUG	2668	CAAGAGACATATTAAGATGAGACAG	3272
287	UGUCUCAUCUUAAUAUGUCUCUUGC	2669	GCAAGAGACATATTAAGATGAGACA	3273
288	GUCUCAUCUUAAUAUGUCUCUUGCU	2670	AGCAAGAGACATATTAAGATGAGAC	3274
289	UCUCAUCUUAAUAUGUCUCUUGCUG	2671	CAGCAAGAGACATATTAAGATGAGA	3275
290	CUCAUCUUAAUAUGUCUCUUGCUGA	2672	TCAGCAAGAGACATATTAAGATGAG	3276
291	UCAUCUUAAUAUGUCUCUUGCUGAU	2673	ATCAGCAAGAGACATATTAAGATGA	3277
292	CAUCUUAAUAUGUCUCUUGCUGAUC	2674	GATCAGCAAGAGACATATTAAGATG	3278
293	AUCUUAAUAUGUCUCUUGCUGAUCU	2675	AGATCAGCAAGAGACATATTAAGAT	3279
294	UCUUAAUAUGUCUCUUGCUGAUCUG	2676	CAGATCAGCAAGAGACATATTAAGA	3280
295	CUUUAAUAUGUCUCUUGCUGAUCUGU	2677	ACAGATCAGCAAGAGACATATTAAG	3281
296	UUAAUAUGUCUCUUGCUGAUCUGUA	2678	TACAGATCAGCAAGAGACATATTAA	3282

Table 30

297	UAAUAUGUCUCUUGCUGAUCUGUAU	2679	ATACAGATCAGCAAGAGACATATTA	3283
298	AAUAUGUCUCUUGCUGAUCUGUAUC	2680	GATACAGATCAGCAAGAGACATATT	3284
299	AUAUGUCUCUUGCUGAUCUGUAUCA	2681	TGATACAGATCAGCAAGAGACATAT	3285
300	UAUGUCUCUUGCUGAUCUGUAUCAU	2682	ATGATACAGATCAGCAAGAGACATA	3286
301	AUGUCUCUUGCUGAUCUGUAUCAUC	2683	GATGATACAGATCAGCAAGAGACAT	3287
302	UGUCUCUUGCUGAUCUGUAUCAUCG	2684	CGATGATACAGATCAGCAAGAGACA	3288
303	GUCUCUUGCUGAUCUGUAUCAUCGU	2685	ACGATGATACAGATCAGCAAGAGAC	3289
304	UCUCUUGCUGAUCUGUAUCAUCGUG	2686	CACGATGATACAGATCAGCAAGAGA	3290
305	CUCUUGCUGAUCUGUAUCAUCGUGA	2687	TCACGATGATACAGATCAGCAAGAG	3291
306	UCUUGCUGAUCUGUAUCAUCGUGAU	2688	ATCACGATGATACAGATCAGCAAGA	3292
307	CUUGCUGAUCUGUAUCAUCGUGAUG	2689	CATCACGATGATACAGATCAGCAAG	3293
308	UUGCUGAUCUGUAUCAUCGUGAUGC	2690	GCATCACGATGATACAGATCAGCAA	3294
309	UGCUGAUCUGUAUCAUCGUGAUGCU	2691	AGCATCACGATGATACAGATCAGCA	3295
310	GCUGAUCUGUAUCAUCGUGAUGCUU	2692	AAGCATCACGATGATACAGATCAGC	3296
311	CUGAUCUGUAUCAUCGUGAUGCUUC	2693	GAAGCATCACGATGATACAGATCAG	3297
312	UGAUCUGUAUCAUCGUGAUGCUUCU	2694	AGAAGCATCACGATGATACAGATCA	3298
313	GAUCUGUAUCAUCGUGAUGCUUCUC	2695	GAGAAGCATCACGATGATACAGATC	3299
314	AUCUGUAUCAUCGUGAUGCUUCUCU	2696	AGAGAAGCATCACGATGATACAGAT	3300
315	UCUGUAUCAUCGUGAUGCUUCUCUG	2697	CAGAGAAGCATCACGATGATACAGA	3301
316	CUGUAUCAUCGUGAUGCUUCUCUGA	2698	TCAGAGAAGCATCACGATGATACAG	3302
317	UGUAUCAUCGUGAUGCUUCUCUGAA	2699	TTACAGAGAAGCATCACGATGATACA	3303
318	GUAUCAUCGUGAUGCUUCUCUGAAG	2700	CTTCAGAGAAGCATCACGATGATAC	3304
319	UAUCAUCGUGAUGCUUCUCUGAAGU	2701	ACTTCAGAGAAGCATCACGATGATA	3305
320	AUCAUCGUGAUGCUUCUCUGAAGUU	2702	AACTTCAGAGAAGCATCACGATGAT	3306
321	UCAUCGUGAUGCUUCUCUGAAGUUC	2703	GAACTTCAGAGAAGCATCACGATGA	3307
322	CAUCGUGAUGCUUCUCUGAAGUUCU	2704	AGAACTTCAGAGAAGCATCACGATG	3308
323	AUCGUGAUGCUUCUCUGAAGUUCUG	2705	CAGAACTTCAGAGAAGCATCACGAT	3309
324	UCGUGAUGCUUCUCUGAAGUUCUGC	2706	GCAGAACTTCAGAGAAGCATCACGA	3310
325	CGUGAUGCUUCUCUGAAGUUCUGCU	2707	AGCAGAACTTCAGAGAAGCATCACG	3311
326	GUGAUGCUUCUCUGAAGUUCUGCUA	2708	TAGCAGAACTTCAGAGAAGCATCAC	3312
327	UGAUGCUUCUCUGAAGUUCUGCUAC	2709	GTAGCAGAACTTCAGAGAAGCATCA	3313
328	GAUGCUUCUCUGAAGUUCUGCUACA	2710	TGTAGCAGAACTTCAGAGAAGCATC	3314
329	AUGCUUCUCUGAAGUUCUGCUACAA	2711	TTGTAGCAGAACTTCAGAGAAGCAT	3315
330	UGCUUCUCUGAAGUUCUGCUACAAC	2712	GTTGTAGCAGAACTTCAGAGAAGCA	3316
331	GCUUCUCUGAAGUUCUGCUACAACC	2713	GGTTGTAGCAGAACTTCAGAGAAGC	3317
332	CUUCUCUGAAGUUCUGCUACAACCU	2714	AGGTTGTAGCAGAACTTCAGAGAAG	3318
333	UUCUCUGAAGUUCUGCUACAACCUC	2715	GAGGTTGTAGCAGAACTTCAGAGAA	3319
334	UCUCUGAAGUUCUGCUACAACCUCU	2716	AGAGGTTGTAGCAGAACTTCAGAGA	3320
335	CUCUGAAGUUCUGCUACAACCUCUA	2717	TAGAGGTTGTAGCAGAACTTCAGAG	3321
336	UCUGAAGUUCUGCUACAACCUCUAG	2718	CTAGAGGTTGTAGCAGAACTTCAGA	3322
337	CUGAAGUUCUGCUACAACCUCUAGA	2719	TCTAGAGGTTGTAGCAGAACTTCAG	3323
338	UGAAGUUCUGCUACAACCUCUAGAU	2720	ATCTAGAGGTTGTAGCAGAACTTCA	3324
339	GAAGUUCUGCUACAACCUCUAGAUC	2721	GATCTAGAGGTTGTAGCAGAACTTC	3325
340	AAGUUCUGCUACAACCUCUAGAUCU	2722	AGATCTAGAGGTTGTAGCAGAACTT	3326
341	AGUUCUGCUACAACCUCUAGAUCUG	2723	CAGATCTAGAGGTTGTAGCAGAACT	3327
342	GUUCUGCUACAACCUCUAGAUCUGC	2724	GCAGATCTAGAGGTTGTAGCAGAAC	3328
343	UUCUGCUACAACCUCUAGAUCUGCA	2725	TGCAGATCTAGAGGTTGTAGCAGAA	3329

Table 30

344	UCUGCUACAACCUCUAGAUCUGCAG	2726	CTGCAGATCTAGAGGTTGTAGCAGA	3330
345	CUGCUACAACCUCUAGAUCUGCAGC	2727	GCTGCAGATCTAGAGGTTGTAGCAG	3331
346	UGCUACAACCUCUAGAUCUGCAGCU	2728	AGCTGCAGATCTAGAGGTTGTAGCA	3332
347	GCUACAACCUCUAGAUCUGCAGCUU	2729	AAGCTGCAGATCTAGAGGTTGTAGC	3333
348	CUACAACCUCUAGAUCUGCAGCUUG	2730	CAAGCTGCAGATCTAGAGGTTGTAG	3334
349	UACAACCUCUAGAUCUGCAGCUUGC	2731	GCAAGCTGCAGATCTAGAGGTTGTA	3335
350	ACAACCUCUAGAUCUGCAGCUUGCC	2732	GGCAAGCTGCAGATCTAGAGGTTGT	3336
351	CAACCUCUAGAUCUGCAGCUUGCCA	2733	TGGCAAGCTGCAGATCTAGAGGTTG	3337
352	AACCUCUAGAUCUGCAGCUUGCCAC	2734	GTGGCAAGCTGCAGATCTAGAGGTT	3338
353	ACCUCUAGAUCUGCAGCUUGCCACA	2735	TGTGGCAAGCTGCAGATCTAGAGGT	3339
354	CCUCUAGAUCUGCAGCUUGCCACAU	2736	ATGTGGCAAGCTGCAGATCTAGAGG	3340
355	CUCUAGAUCUGCAGCUUGCCACAUC	2737	GATGTGGCAAGCTGCAGATCTAGAG	3341
356	UCUAGAUCUGCAGCUUGCCACAUCA	2738	TGATGTGGCAAGCTGCAGATCTAGA	3342
357	CUAGAUCUGCAGCUUGCCACAUCAG	2739	CTGATGTGGCAAGCTGCAGATCTAG	3343
358	UAGAUCUGCAGCUUGCCACAUCAGC	2740	GCTGATGTGGCAAGCTGCAGATCTA	3344
368	GCUUGCCACAUCAGCUUAAAUCUG	2741	CAGATTTTAAGCTGATGTGGCAAGC	3345
369	CUUGCCACAUCAGCUUAAAUCUGU	2742	ACAGATTTTAAGCTGATGTGGCAAG	3346
370	UUGCCACAUCAGCUUAAAUCUGUC	2743	GACAGATTTTAAGCTGATGTGGCAA	3347
371	UGCCACAUCAGCUUAAAUCUGUCA	2744	TGACAGATTTTAAGCTGATGTGGCA	3348
372	GCCACAUCAGCUUAAAUCUGUCAU	2745	ATGACAGATTTTAAGCTGATGTGGC	3349
373	CCACAUCAGCUUAAAUCUGUCAUC	2746	GATGACAGATTTTAAGCTGATGTGG	3350
374	CACAUCAGCUUAAAUCUGUCAUCC	2747	GGATGACAGATTTTAAGCTGATGTG	3351
375	ACAUCAGCUUAAAUCUGUCAUCCC	2748	GGGATGACAGATTTTAAGCTGATGT	3352
376	CAUCAGCUUAAAUCUGUCAUCCCA	2749	TGGATGACAGATTTTAAGCTGATG	3353
377	AUCAGCUUAAAUCUGUCAUCCCAU	2750	ATGGGATGACAGATTTTAAGCTGAT	3354
378	UCAGCUUAAAUCUGUCAUCCCAUG	2751	CATGGGATGACAGATTTTAAGCTGA	3355
379	CAGCUUAAAUCUGUCAUCCCAUGC	2752	GCATGGGATGACAGATTTTAAGCTG	3356
380	AGCUUAAAUCUGUCAUCCCAUGCA	2753	TGCATGGGATGACAGATTTTAAGCT	3357
381	GCUUAAAUCUGUCAUCCCAUGCAG	2754	CTGCATGGGATGACAGATTTTAAGC	3358
382	CUUAAAUCUGUCAUCCCAUGCAGA	2755	TCTGCATGGGATGACAGATTTTAAG	3359
383	UUAAAUCUGUCAUCCCAUGCAGAC	2756	GTCTGCATGGGATGACAGATTTTAA	3360
384	UAAAUCUGUCAUCCCAUGCAGACA	2757	TGTCTGCATGGGATGACAGATTTTA	3361
391	UGUCAUCCCAUGCAGACAGGAAAAC	2758	GTTTTCCTGTCTGCATGGGATGACA	3362
392	GUCAUCCCAUGCAGACAGGAAAACA	2759	TGTTTTCCTGTCTGCATGGGATGAC	3363
393	UCAUCCCAUGCAGACAGGAAAACAA	2760	TTGTTTTCCTGTCTGCATGGGATGA	3364
394	CAUCCCAUGCAGACAGGAAAACAAU	2761	ATTGTTTTCCTGTCTGCATGGGATG	3365
395	AUCCCAUGCAGACAGGAAAACAAUA	2762	TATGTTTTCCTGTCTGCATGGGAT	3366
396	UCCCAUGCAGACAGGAAAACAAUUA	2763	ATATTGTTTTCCTGTCTGCATGGGA	3367
397	CCAUGCAGACAGGAAAACAAUAUU	2764	AATATTGTTTTCCTGTCTGCATGGG	3368
398	CCAUGCAGACAGGAAAACAAUAUUG	2765	CAATATTGTTTTCCTGTCTGCATGG	3369
399	CAUGCAGACAGGAAAACAAUAUUGU	2766	ACAATATTGTTTTCCTGTCTGCATG	3370
400	AUGCAGACAGGAAAACAAUAUUGUA	2767	TACAATATTGTTTTCCTGTCTGCAT	3371
401	UGCAGACAGGAAAACAAUAUUGUAU	2768	ATACAATATTGTTTTCCTGTCTGCA	3372
426	AACAGACCACUCCUGAGUAGAAGA	2769	TCTTCTACTCAGGAAGTGGTCTGTT	3373
427	ACAGACCACUCCUGAGUAGAAGAG	2770	CTCTTCTACTCAGGAAGTGGTCTGT	3374
428	CAGACCACUCCUGAGUAGAAGAGU	2771	ACTCTTCTACTCAGGAAGTGGTCTG	3375
430	GACCACUCCUGAGUAGAAGAGUUU	2772	AAACTCTTCTACTCAGGAAGTGGTC	3376

Table 30

431	ACCACUCCUGAGUAGAAGAGUUUC	2773	GAAACTCTTCTACTCAGGAAGTGGT	3377
432	CCACUCCUGAGUAGAAGAGUUUCU	2774	AGAAACTCTTCTACTCAGGAAGTGG	3378
445	AGAAGAGUUUCUUUGUGAAAAGGUC	2775	GACCTTTTCACAAAGAACTCTTCT	3379
446	GAAGAGUUUCUUUGUGAAAAGGUCA	2776	TGACCTTTTCACAAAGAACTCTTC	3380
447	AAGAGUUUCUUUGUGAAAAGGUCAA	2777	TTGACCTTTTCACAAAGAACTCTT	3381
448	AGAGUUUCUUUGUGAAAAGGUCAAG	2778	CTTGACCTTTTCACAAAGAACTCT	3382
449	GAGUUUCUUUGUGAAAAGGUCAAGA	2779	TCTTGACCTTTTCACAAAGAACTC	3383
450	AGUUUCUUUGUGAAAAGGUCAAGAU	2780	ATCTTGACCTTTTCACAAAGAACT	3384
451	GUUUCUUUGUGAAAAGGUCAAGAUU	2781	AATCTTGACCTTTTCACAAAGAAAC	3385
452	UUUCUUUGUGAAAAGGUCAAGAUUA	2782	TAATCTTGACCTTTTCACAAAGAAA	3386
453	UUCUUUGUGAAAAGGUCAAGAUUAA	2783	TTAATCTTGACCTTTTCACAAAGAA	3387
504	AUUCUUCUGUUGGAUCUUGUAAACA	2784	TGTTTACAAGATCCAACAGATGAAT	3388
505	UUCUUCUGUUGGAUCUUGUAAACAU	2785	ATGTTTACAAGATCCAACAGATGAA	3389
506	UCAUCUGUUGGAUCUUGUAAACAUG	2786	CATGTTTACAAGATCCAACAGATGA	3390
507	CAUCUGUUGGAUCUUGUAAACAUGA	2787	TCATGTTTACAAGATCCAACAGATG	3391
508	AUCUGUUGGAUCUUGUAAACAUGAA	2788	TTCATGTTTACAAGATCCAACAGAT	3392
509	UCUGUUGGAUCUUGUAAACAUGAAA	2789	TTTCATGTTTACAAGATCCAACAGA	3393
510	CUGUUGGAUCUUGUAAACAUGAAAA	2790	TTTTCATGTTTACAAGATCCAACAG	3394
511	UGUUGGAUCUUGUAAACAUGAAAAG	2791	CTTTTCATGTTTACAAGATCCAACA	3395
512	GUUGGAUCUUGUAAACAUGAAAAGG	2792	CCTTTTCATGTTTACAAGATCCAAC	3396
513	UUGGAUCUUGUAAACAUGAAAAGGG	2793	CCCTTTTCATGTTTACAAGATCCAA	3397
514	UGGAUCUUGUAAACAUGAAAAGGGC	2794	GCCCTTTTCATGTTTACAAGATCCA	3398
515	GGAUCUUGUAAACAUGAAAAGGGCU	2795	AGCCCTTTTCATGTTTACAAGATCC	3399
516	GAUCUUGUAAACAUGAAAAGGGCUU	2796	AAGCCCTTTTCATGTTTACAAGATC	3400
517	AUCUUGUAAACAUGAAAAGGGCUUU	2797	AAAGCCCTTTTCATGTTTACAAGAT	3401
518	UCUUGUAAACAUGAAAAGGGCUUUA	2798	TAAAGCCCTTTTCATGTTTACAAGA	3402
519	CUUGUAAACAUGAAAAGGGCUUUAU	2799	ATAAGCCCTTTTCATGTTTACAAG	3403
520	UUGUAAACAUGAAAAGGGCUUUAUU	2800	AATAAGCCCTTTTCATGTTTACAA	3404
521	UGUAAACAUGAAAAGGGCUUUAUUU	2801	AAATAAGCCCTTTTCATGTTTACA	3405
522	GUAAACAUGAAAAGGGCUUUAUUUU	2802	AAAATAAGCCCTTTTCATGTTTAC	3406
531	AAAAGGGCUUUAUUUCAAUUUA	2803	TAATTTTGAATAAAGCCCTTTT	3407
532	AAAGGGCUUUAUUUCAAUUUAUA	2804	TTAATTTTGAATAAAGCCCTTTT	3408
533	AAGGGCUUUAUUUCAAUUUAUAC	2805	GTTAATTTTGAATAAAGCCCTT	3409
534	AGGGCUUUAUUUCAAUUUAUACU	2806	AGTTAATTTTGAATAAAGCCCT	3410
535	GGGCUUUAUUUCAAUUUAUACUU	2807	AAGTTAATTTTGAATAAAGCCC	3411
570	GUUAAAAUGCAACUGUUGAUUUC	2808	GGAAATCAACAGTTGCATTTTATAC	3412
571	UAUAAAAUGCAACUGUUGAUUUCU	2809	AGGAAATCAACAGTTGCATTTTATA	3413
572	AUAAAAUGCAACUGUUGAUUUCUC	2810	GAGGAAATCAACAGTTGCATTTTAT	3414
573	UAAAAUGCAACUGUUGAUUUCUCA	2811	TGAGGAAATCAACAGTTGCATTTTA	3415
574	AAAAUGCAACUGUUGAUUUCUCA	2812	TTGAGGAAATCAACAGTTGCATTTT	3416
586	UGAUUUCUCAACAUGGCUCACAA	2813	TTGTGAGCCATGTTGAGGAAATCAA	3417
587	UGAUUUCUCAACAUGGCUCACAAA	2814	TTGTGAGCCATGTTGAGGAAATCA	3418
588	GAUUUUCUCAACAUGGCUCACAAU	2815	ATTTGTGAGCCATGTTGAGGAAATC	3419
589	AUUUCCUCAACAUGGCUCACAAUU	2816	AATTTGTGAGCCATGTTGAGGAAAT	3420
590	UUUCCUCAACAUGGCUCACAAUUU	2817	AAATTTGTGAGCCATGTTGAGGAAA	3421
591	UUCUCAACAUGGCUCACAAUUUC	2818	GAAATTTGTGAGCCATGTTGAGGAA	3422
592	UCCUCAACAUGGCUCACAAUUUCU	2819	AGAAATTTGTGAGCCATGTTGAGGA	3423

Table 30

593	CCUCAACAUGGCUCACAAAUUUCUA	2820	TAGAAATTTGTGAGCCATGTTGAGG	3424
594	CUCAACAUGGCUCACAAAUUUCUAU	2821	ATAGAAATTTGTGAGCCATGTTGAG	3425
595	UCAACAUGGCUCACAAAUUUCUAUC	2822	GATAGAAATTTGTGAGCCATGTTGA	3426
596	CAACAUGGCUCACAAAUUUCUAUCC	2823	GGATAGAAATTTGTGAGCCATGTTG	3427
597	AACAUGGCUCACAAAUUUCUAUCCC	2824	GGGATAGAAATTTGTGAGCCATGTT	3428
598	ACAUGGCUCACAAAUUUCUAUCCCA	2825	TGGGATAGAAATTTGTGAGCCATGT	3429
599	CAUGGCUCACAAAUUUCUAUCCCAA	2826	TTGGGATAGAAATTTGTGAGCCATG	3430
600	AUGGCUCACAAAUUUCUAUCCCAAA	2827	TTTGGGATAGAAATTTGTGAGCCAT	3431
601	UGGCUCACAAAUUUCUAUCCCAAAU	2828	ATTTGGGATAGAAATTTGTGAGCCA	3432
602	GGCUCACAAAUUUCUAUCCCAAAUC	2829	GATTTGGGATAGAAATTTGTGAGCC	3433
603	GCUCACAAAUUUCUAUCCCAAAUCU	2830	AGATTTGGGATAGAAATTTGTGAGC	3434
604	CUCACAAAUUUCUAUCCCAAAUCUU	2831	AAGATTTGGGATAGAAATTTGTGAG	3435
605	UCACAAAUUUCUAUCCCAAAUCUUU	2832	AAAGATTTGGGATAGAAATTTGTGA	3436
606	CACAAAUUUCUAUCCCAAAUCUUUU	2833	AAAAGATTTGGGATAGAAATTTGTG	3437
607	ACAAAUUUCUAUCCCAAAUCUUUUC	2834	GAAAAGATTTGGGATAGAAATTTGT	3438
608	CAAAUUCUAUCCCAAAUCUUUUCU	2835	AGAAAAGATTTGGGATAGAAATTTG	3439
609	AAAUUUCUAUCCCAAAUCUUUUCUG	2836	CAGAAAAGATTTGGGATAGAAATTT	3440
610	AAUUCUAUCCCAAAUCUUUUCUGA	2837	TCAGAAAAGATTTGGGATAGAAATT	3441
611	AUUCUAUCCCAAAUCUUUUCUGAA	2838	TTCAGAAAAGATTTGGGATAGAAAT	3442
612	UUUCUAUCCCAAAUCUUUUCUGAAG	2839	CTTCAGAAAAGATTTGGGATAGAAA	3443
613	UUCUAUCCCAAAUCUUUUCUGAAGA	2840	TCTTCAGAAAAGATTTGGGATAGAA	3444
644	GUUUAGUUUUAAAACUGCACUGCCA	2841	TGGCAGTGCAGTTTTAAACTAAAC	3445
645	UUUAGUUUUAAAACUGCACUGCCAA	2842	TTGGCAGTGCAGTTTTAAACTAAA	3446
646	UUAGUUUUAAAACUGCACUGCCAAC	2843	GTTGGCAGTGCAGTTTTAAACTAA	3447
647	UAGUUUUAAAACUGCACUGCCAACA	2844	TGTTGGCAGTGCAGTTTTAAACTA	3448
648	AGUUUUAAAACUGCACUGCCAACAA	2845	TTGTTGGCAGTGCAGTTTTAAACT	3449
649	GUUUUAAAACUGCACUGCCAACAAG	2846	CTTGTTGGCAGTGCAGTTTTAAAC	3450
650	UUUUAAAACUGCACUGCCAACAAGU	2847	ACTTGTGGCAGTGCAGTTTTAAAA	3451
651	UUUAAAACUGCACUGCCAACAAGUU	2848	AACTTGTGGCAGTGCAGTTTTAAA	3452
652	UUAAAACUGCACUGCCAACAAGUUC	2849	GAACTTGTGGCAGTGCAGTTTTAA	3453
653	UAAAACUGCACUGCCAACAAGUUCA	2850	TGAACTTGTGGCAGTGCAGTTTTA	3454
654	AAAACUGCACUGCCAACAAGUUCAC	2851	GTGAACTTGTGGCAGTGCAGTTTT	3455
655	AAACUGCACUGCCAACAAGUUCACU	2852	AGTGAACCTGTTGGCAGTGCAGTTT	3456
656	AACUGCACUGCCAACAAGUUCACUU	2853	AAGTGAACCTGTTGGCAGTGCAGTT	3457
657	ACUGCACUGCCAACAAGUUCACUUC	2854	GAAGTGAACCTGTTGGCAGTGCAGT	3458
658	CUGCACUGCCAACAAGUUCACUUCA	2855	TGAAGTGAACCTGTTGGCAGTGCAG	3459
659	UGCACUGCCAACAAGUUCACUUCAU	2856	ATGAAGTGAACCTGTTGGCAGTGCA	3460
660	GCACUGCCAACAAGUUCACUUCAUA	2857	TATGAAGTGAACCTGTTGGCAGTGC	3461
661	CACUGCCAACAAGUUCACUUCAUAU	2858	ATATGAAGTGAACCTGTTGGCAGTG	3462
662	ACUGCCAACAAGUUCACUUCAUAUA	2859	TATATGAAGTGAACCTGTTGGCAGT	3463
663	CUGCCAACAAGUUCACUUCAUAUAU	2860	ATATATGAAGTGAACCTGTTGGCAG	3464
755	UAAGUAUUUUUCAGGUCUUCACCAA	2861	TTGGTGAAGACCTGAAAAATACTTA	3465
756	AAGUAUUUUUCAGGUCUUCACCAAG	2862	CTTGGTGAAGACCTGAAAAATACTT	3466
757	AGUAUUUUUCAGGUCUUCACCAAGU	2863	ACTTGGTGAAGACCTGAAAAATACT	3467
760	AUUUUUCAGGUCUUCACCAAGUAUC	2864	GATACTTGGTGAAGACCTGAAAAAT	3468
761	UUUUUCAGGUCUUCACCAAGUAUCA	2865	TGATACTTGGTGAAGACCTGAAAAA	3469
762	UUUUCAGGUCUUCACCAAGUAUCAA	2866	TTGATACTTGGTGAAGACCTGAAAA	3470

Table 30

763	UUUCAGGUCUUCACCAAGUAUCAA	2867	TTTGATACTTGGTGAAGACCTGAAA	3471
764	UUCAGGUCUUCACCAAGUAUCAAAG	2868	CTTTGATACTTGGTGAAGACCTGAA	3472
765	UCAGGUCUUCACCAAGUAUCAAAGU	2869	ACTTTGATACTTGGTGAAGACCTGA	3473
766	CAGGUCUUCACCAAGUAUCAAAGUA	2870	TACTTTGATACTTGGTGAAGACCTG	3474
813	AUUCAAAAUAGUCCACUGACUCCUC	2871	GAGGAGTCAGTGGACTATTTTGAAT	3475
814	UUCAAAAUAGUCCACUGACUCCUCA	2872	TGAGGAGTCAGTGGACTATTTTGAA	3476
815	UCAAAAAUAGUCCACUGACUCCUCAC	2873	GTGAGGAGTCAGTGGACTATTTTGA	3477
816	CAAAAAUAGUCCACUGACUCCUCACA	2874	TGTGAGGAGTCAGTGGACTATTTTG	3478
817	AAAAUAGUCCACUGACUCCUCACAU	2875	ATGTGAGGAGTCAGTGGACTATTTT	3479
818	AAAUAGUCCACUGACUCCUCACAUC	2876	GATGTGAGGAGTCAGTGGACTATTT	3480
819	AAUAGUCCACUGACUCCUCACAUCU	2877	AGATGTGAGGAGTCAGTGGACTATT	3481
820	AUAGUCCACUGACUCCUCACAUCUG	2878	CAGATGTGAGGAGTCAGTGGACTAT	3482
821	UAGUCCACUGACUCCUCACAUCUGU	2879	ACAGATGTGAGGAGTCAGTGGACTA	3483
822	AGUCCACUGACUCCUCACAUCUGUU	2880	AACAGATGTGAGGAGTCAGTGGACT	3484
823	GUCCACUGACUCCUCACAUCUGUUA	2881	TAACAGATGTGAGGAGTCAGTGGAC	3485
824	UCCACUGACUCCUCACAUCUGUUUAU	2882	ATAACAGATGTGAGGAGTCAGTGGGA	3486
825	CCACUGACUCCUCACAUCUGUUUAUC	2883	GATAACAGATGTGAGGAGTCAGTGG	3487
911	UUUUUCUAUGCCACAUUAACAUCUU	2884	AAGATGTTAATGTGGCATAGAAAAA	3488
912	UUUUCUAUGCCACAUUAACAUCUUU	2885	AAAGATGTTAATGTGGCATAGAAAA	3489
913	UUUCUAUGCCACAUUAACAUCUUUU	2886	AAAAGATGTTAATGTGGCATAGAAA	3490
919	UGCCACAUUAACAUCUUUUAAGUU	2887	AACTTTAAAAGATGTTAATGTGGCA	3491
920	GCCACAUUAACAUCUUUUAAGUUG	2888	CAACTTTAAAAGATGTTAATGTGGC	3492
948	AGAAUCAAGUAUGGAAAAGUAAGGC	2889	GCCTTACTTTTCCATACTTGATTCT	3493
949	GAAUCAAGUAUGGAAAAGUAAGGCC	2890	GGCCTTACTTTTCCATACTTGATTCT	3494
950	AAUCAAGUAUGGAAAAGUAAGGCCA	2891	TGGCCTTACTTTTCCATACTTGATT	3495
959	UGGAAAAGUAAGGCCAUACUCUUAC	2892	GTAAGAGTATGGCCTTACTTTTCCA	3496
960	GGAAAAGUAAGGCCAUACUCUUACA	2893	TGTAAGAGTATGGCCTTACTTTTCC	3497
1067	CAUAUGAUCAACAGAUGAGAACUGG	2894	CCAGTTCTCATCTGTTGATCATATG	3498
1069	UAUGAUCAACAGAUGAGAACUGGUG	2895	CACCAGTTCTCATCTGTTGATCATA	3499
1070	AUGAUCAACAGAUGAGAACUGGUGG	2896	CCACCAGTTCTCATCTGTTGATCAT	3500
1071	UGAUCAACAGAUGAGAACUGGUGGU	2897	ACCACCAGTTCTCATCTGTTGATCA	3501
1072	GAUCAACAGAUGAGAACUGGUGGUU	2898	AACCACCAGTTCTCATCTGTTGATC	3502
1073	AUCAACAGAUGAGAACUGGUGGUUA	2899	TAACCACCAGTTCTCATCTGTTGAT	3503
1074	UCAACAGAUGAGAACUGGUGGUUAA	2900	TTAACCACCAGTTCTCATCTGTTGA	3504
1075	CAACAGAUGAGAACUGGUGGUUAAU	2901	ATTAACCACCAGTTCTCATCTGTTG	3505
1078	CAGAUGAGAACUGGUGGUUAAUUG	2902	CATATTAAACCACCAGTTCTCATCTG	3506
1080	GAUGAGAACUGGUGGUUAAUUGUG	2903	CACATATTAAACCACCAGTTCTCATC	3507
1081	AUGAGAACUGGUGGUUAAUUGUGA	2904	TCACATATTAAACCACCAGTTCTCAT	3508
1082	UGAGAACUGGUGGUUAAUUGUGAC	2905	GTCACATATTAAACCACCAGTTCTCA	3509
1083	GAGAACUGGUGGUUAAUUGUGACA	2906	TGTCACATATTAAACCACCAGTTCTC	3510
1086	AACUGGUGGUUAAUUGUGACAGUG	2907	CACTGTCACATATTAAACCACCAGTT	3511
1087	ACUGGUGGUUAAUUGUGACAGUGA	2908	TCACTGTCACATATTAAACCACCAGT	3512
1088	CUGGUGGUUAAUUGUGACAGUGAG	2909	CTCACTGTCACATATTAAACCACCAG	3513
1089	UGGUGGUUAAUUGUGACAGUGAGA	2910	TCTCACTGTCACATATTAAACCACCA	3514
1141	CAGAAUCUAAUCUUAUUAAGGCA	2911	TGCCTTAAATGAAGATTAGATTCTG	3515
1150	AUCUUAUUAAGGCACUGUAGUGA	2912	TCACTACAGTGCCTTAAATGAAGAT	3516
1151	UCUUAUUAAGGCACUGUAGUGAA	2913	TTCACTACAGTGCCTTAAATGAAGA	3517

Table 30

1153	UUCAUUUAAGGCACUGUAGUGAAUU	2914	AATCACTACAGTGCCTTAAATGAA	3518
1161	AGGCACUGUAGUGAAUUUUCUGAGC	2915	GCTCAGATAATTCACAGTGCCT	3519
1162	GGCACUGUAGUGAAUUUUCUGAGCU	2916	AGCTCAGATAATTCACAGTGCC	3520
1211	UAUCUUUGGAAUCAUGAAACCUUAA	2917	TTAAGGTTTCATGATTCCAAAGATA	3521
1212	AUCUUUGGAAUCAUGAAACCUUAG	2918	CTTAAGGTTTCATGATTCCAAAGAT	3522
1213	UCUUUGGAAUCAUGAAACCUUAGA	2919	TCTTAAGGTTTCATGATTCCAAAGA	3523
1214	CUUUGGAAUCAUGAAACCUUAGAC	2920	GTCTTAAGGTTTCATGATTCCAAAG	3524
1215	UUUGGAAUCAUGAAACCUUAGACU	2921	AGTCTTAAGGTTTCATGATTCCAAA	3525
1216	UUGGAAUCAUGAAACCUUAGACUU	2922	AAGTCTTAAGGTTTCATGATTCCAA	3526
1217	UGGAAUCAUGAAACCUUAGACUUC	2923	GAAGTCTTAAGGTTTCATGATTCCA	3527
1218	GGAAUCAUGAAACCUUAGACUUCA	2924	TGAAGTCTTAAGGTTTCATGATTCC	3528
1223	CAUGAAACCUUAGACUUCAGAAUG	2925	CATTCTGAAGTCTTAAGGTTTCATG	3529
1230	CCUUAAGACUUCAGAAUGAUUUUGC	2926	GCAAAATCATTCTGAAGTCTTAAGG	3530
1231	CUUAAGACUUCAGAAUGAUUUUGCA	2927	TGCAAAATCATTCTGAAGTCTTAAG	3531
1232	UUAAGACUUCAGAAUGAUUUUGCAG	2928	CTGCAAAATCATTCTGAAGTCTTAA	3532
1233	UAAGACUUCAGAAUGAUUUUGCAGG	2929	CCTGCAAAATCATTCTGAAGTCTTA	3533
1234	AAGACUUCAGAAUGAUUUUGCAGGU	2930	ACCTGCAAAATCATTCTGAAGTCTT	3534
1235	AGACUUCAGAAUGAUUUUGCAGGUU	2931	AACCTGCAAAATCATTCTGAAGTCT	3535
1236	GACUUCAGAAUGAUUUUGCAGGUUG	2932	CAACCTGCAAAATCATTCTGAAGTC	3536
1237	ACUUCAGAAUGAUUUUGCAGGUUGU	2933	ACAACCTGCAAAATCATTCTGAAGT	3537
1238	CUUCAGAAUGAUUUUGCAGGUUGUC	2934	GACAACCTGCAAAATCATTCTGAAG	3538
1239	UUCAGAAUGAUUUUGCAGGUUGUCU	2935	AGACAACCTGCAAAATCATTCTGAA	3539
1240	UCAGAAUGAUUUUGCAGGUUGUCUU	2936	AAGACAACCTGCAAAATCATTCTGA	3540
1241	CAGAAUGAUUUUGCAGGUUGUCUUC	2937	GAAGACAACCTGCAAAATCATTCTG	3541
1242	AGAAUGAUUUUGCAGGUUGUCUUC	2938	GGAAGACAACCTGCAAAATCATTCT	3542
1243	GAAUGAUUUUGCAGGUUGUCUCCA	2939	TGGAAGACAACCTGCAAAATCATT	3543
1244	AAUGAUUUUGCAGGUUGUCUCCA	2940	ATGGAAGACAACCTGCAAAATCATT	3544
1245	AUGAUUUUGCAGGUUGUCUCCA	2941	AATGGAAGACAACCTGCAAAATCAT	3545
1246	UGAUUUUGCAGGUUGUCUCCA	2942	GAATGGAAGACAACCTGCAAAATCA	3546
1247	GAUUUGCAGGUUGUCUCCA	2943	GGAATGGAAGACAACCTGCAAAATC	3547
1248	AUUUGCAGGUUGUCUCCA	2944	TGGAATGGAAGACAACCTGCAAAAT	3548
1249	UUUGCAGGUUGUCUCCA	2945	CTGGAATGGAAGACAACCTGCAAAA	3549
1250	UUUGCAGGUUGUCUCCA	2946	GCTGGAATGGAAGACAACCTGCAAA	3550
1251	UUGCAGGUUGUCUCCA	2947	GGCTGGAATGGAAGACAACCTGCAA	3551
1252	UGCAGGUUGUCUCCA	2948	AGGCTGGAATGGAAGACAACCTGCA	3552
1253	GCAGGUUGUCUCCA	2949	TAGGCTGGAATGGAAGACAACCTGC	3553
1254	CAGGUUGUCUCCA	2950	TTAGGCTGGAATGGAAGACAACCTG	3554
1255	AGGUUGUCUCCA	2951	GTTAGGCTGGAATGGAAGACAACCT	3555
1256	GGUUGUCUCCA	2952	TGTTAGGCTGGAATGGAAGACAACC	3556
1257	GUUGUCUCCA	2953	ATGTTAGGCTGGAATGGAAGACAAC	3557
1258	UUGUCUCCA	2954	GATGTTAGGCTGGAATGGAAGACA	3558
1259	UGUCUCCA	2955	GGATGTTAGGCTGGAATGGAAGACA	3559
1260	GUCUCCA	2956	TGGATGTTAGGCTGGAATGGAAGAC	3560
1261	UCUCCA	2957	TTGGATGTTAGGCTGGAATGGAAGA	3561
1262	CUCCA	2958	ATTGGATGTTAGGCTGGAATGGAAG	3562
1263	UCCA	2959	CATTGGATGTTAGGCTGGAATGGAA	3563
1264	UCCA	2960	GCATTGGATGTTAGGCTGGAATGGA	3564

Table 30

1265	CCAUCCAGCCUAACAUCCAAUGCA	2961	TGCATTGGATGTTAGGCTGGAATGG	3565
1266	CAUCCAGCCUAACAUCCAAUGCAG	2962	CTGCATTGGATGTTAGGCTGGAATG	3566
1267	AUCCAGCCUAACAUCCAAUGCAGG	2963	CCTGCATTGGATGTTAGGCTGGAAT	3567
1274	CCUAAUCAUCCAAUGCAGGCAAGGAA	2964	TTCCTTGCCTGCATTGGATGTTAGG	3568
1275	CUAAUCAUCCAAUGCAGGCAAGGAAA	2965	TTTCCTTGCCTGCATTGGATGTTAG	3569
1276	UAACAUCCAAUGCAGGCAAGGAAAA	2966	TTTCCTTGCCTGCATTGGATGTTA	3570
1277	AACAUCCAAUGCAGGCAAGGAAAAU	2967	ATTTTCCTTGCCTGCATTGGATGTT	3571
1278	ACAUCCAAUGCAGGCAAGGAAAAUA	2968	TATTTTCCTTGCCTGCATTGGATGT	3572
1279	CAUCCAAUGCAGGCAAGGAAAAUAA	2969	TTATTTTCCTTGCCTGCATTGGATG	3573
1280	AUCCAAUGCAGGCAAGGAAAAUAAA	2970	TTTATTTTCCTTGCCTGCATTGGAT	3574
1281	UCCAAUGCAGGCAAGGAAAAUAAAA	2971	TTTTATTTTCCTTGCCTGCATTGGA	3575
1282	CCAAUGCAGGCAAGGAAAAUAAAAG	2972	CTTTTATTTTCCTTGCCTGCATTGG	3576
1283	CAAUGCAGGCAAGGAAAAUAAAAGA	2973	TCTTTTATTTTCCTTGCCTGCATTG	3577
1284	AAUGCAGGCAAGGAAAAUAAAAGAU	2974	ATCTTTTATTTTCCTTGCCTGCATT	3578
1285	AUGCAGGCAAGGAAAAUAAAAGAUU	2975	AATCTTTTATTTTCCTTGCCTGCAT	3579
1286	UGCAGGCAAGGAAAAUAAAAGAUUU	2976	AAATCTTTTATTTTCCTTGCCTGCA	3580
1287	GCAGGCAAGGAAAAUAAAAGAUUUC	2977	GAAATCTTTTATTTTCCTTGCCTGC	3581
1301	UAAAAGAUUCCAGUGACAGAAAAA	2978	TTTTTCTGTCACTGGAAATCTTTTA	3582
1302	AAAAGAUUCCAGUGACAGAAAAAU	2979	ATTTTCTGTCACTGGAAATCTTTT	3583
1393	UAUAUUUUGAAAUAGAACUUGUUGGC	2980	GCCAACAAGTTCATTTCAAAATATA	3584
1394	AUAUUUUGAAAUAGAACUUGUUGGCC	2981	GGCCAACAAGTTCATTTCAAAATAT	3585
1395	UAUUUUGAAAUAGAACUUGUUGGCC	2982	GGGCAACAAGTTCATTTCAAAATA	3586
1396	AUUUUGAAAUAGAACUUGUUGGCCCA	2983	TGGGCAACAAGTTCATTTCAAAAT	3587
1397	UUUUGAAAUAGAACUUGUUGGCCCAU	2984	ATGGGCAACAAGTTCATTTCAAA	3588
1398	UUUGAAAUAGAACUUGUUGGCCCAUC	2985	GATGGGCAACAAGTTCATTTCAAA	3589
1399	UUGAAAUAGAACUUGUUGGCCCAUCU	2986	AGATGGGCAACAAGTTCATTTCAA	3590
1400	UGAAAUAGAACUUGUUGGCCCAUCUA	2987	TAGATGGGCAACAAGTTCATTTC	3591
1401	GAAUAGAACUUGUUGGCCCAUCUAU	2988	ATAGATGGGCAACAAGTTCATTTC	3592
1402	AAUAGAACUUGUUGGCCCAUCUAUU	2989	AATAGATGGGCAACAAGTTCATT	3593
1403	AAUGAACUUGUUGGCCCAUCUAUUA	2990	TAATAGATGGGCAACAAGTTCATT	3594
1404	AUGAACUUGUUGGCCCAUCUAUUAC	2991	GTAATAGATGGGCAACAAGTTCAT	3595
1405	UGAACUUGUUGGCCCAUCUAUUACA	2992	TGTAATAGATGGGCAACAAGTTC	3596
1406	GAACUUGUUGGCCCAUCUAUUACAU	2993	ATGTAATAGATGGGCAACAAGTTC	3597
1407	AACUUGUUGGCCCAUCUAUUACAUC	2994	GATGTAATAGATGGGCAACAAGTT	3598
1408	ACUUGUUGGCCCAUCUAUUACAUCU	2995	AGATGTAATAGATGGGCAACAAGT	3599
1409	CUUGUUGGCCCAUCUAUUACAUCUA	2996	TAGATGTAATAGATGGGCAACAAG	3600
1410	UUGUUGGCCCAUCUAUUACAUCUAC	2997	GTAGATGTAATAGATGGGCAACAA	3601
1411	UGUUGGCCCAUCUAUUACAUCUACA	2998	TGTAGATGTAATAGATGGGCAACA	3602
1412	GUUGGCCCAUCUAUUACAUCUACAG	2999	CTGTAGATGTAATAGATGGGCAAC	3603
1413	UUGGCCCAUCUAUUACAUCUACAGC	3000	GCTGTAGATGTAATAGATGGGCA	3604
1414	UGGCCCAUCUAUUACAUCUACAGCU	3001	AGCTGTAGATGTAATAGATGGGCC	3605
1415	GGGCCAUCUAUUACAUCUACAGCUG	3002	CAGCTGTAGATGTAATAGATGGGCC	3606
1416	GCCCAUCUAUUACAUCUACAGCUGA	3003	TCAGCTGTAGATGTAATAGATGGGC	3607
1422	CUAUACAUCUACAGCUGACCCUUG	3004	CAAGGGTCAGCTGTAGATGTAATAG	3608
1423	UAUACAUCUACAGCUGACCCUUGA	3005	TCAAGGGTCAGCTGTAGATGTAATA	3609
1424	AUUACAUCUACAGCUGACCCUUGAA	3006	TTCAAGGGTCAGCTGTAGATGTAAT	3610
1425	UUACAUCUACAGCUGACCCUUGAAC	3007	GTTCAAGGGTCAGCTGTAGATGTAA	3611

Table 30

1426	UACAUCUACAGCUGACCCUUGAACA	3008	TGTTCAAGGGTCAGCTGTAGATGTA	3612
1427	ACAUCUACAGCUGACCCUUGAACA	3009	ATGTTCAAGGGTCAGCTGTAGATGT	3613
1428	CAUCUACAGCUGACCCUUGAACAUG	3010	CATGTTCAAGGGTCAGCTGTAGATG	3614
1429	AUCUACAGCUGACCCUUGAACAUGG	3011	CCATGTTCAAGGGTCAGCTGTAGAT	3615
1442	CCUUGAACAUGGGGUUAGGGGAGC	3012	GCTCCCCTAACCCCATGTTCAAGG	3616
1443	CUUGAACAUGGGGUUAGGGGAGCU	3013	AGCTCCCCTAACCCCATGTTCAAG	3617
1444	UUGAACAUGGGGUUAGGGGAGCUG	3014	CAGCTCCCCTAACCCCATGTTCAA	3618
1445	UGAACAUGGGGUUAGGGGAGCUGA	3015	TCAGCTCCCCTAACCCCATGTTCA	3619
1446	GAACAUGGGGUUAGGGGAGCUGAC	3016	GTCAGCTCCCCTAACCCCATGTTT	3620
1447	AACAUGGGGUUAGGGGAGCUGACA	3017	TGTCAGCTCCCCTAACCCCATGTT	3621
1448	ACAUGGGGUUAGGGGAGCUGACAA	3018	TTGTCAGCTCCCCTAACCCCATGT	3622
1449	CAUGGGGUUAGGGGAGCUGACAAU	3019	ATTGTCAGCTCCCCTAACCCCATG	3623
1450	AUGGGGUUAGGGGAGCUGACAAUU	3020	AATTGTCAGCTCCCCTAACCCCAT	3624
1451	UGGGGUUAGGGGAGCUGACAAUUC	3021	GAATTGTCAGCTCCCCTAACCCCA	3625
1452	GGGGGUUAGGGGAGCUGACAAUUCG	3022	CGAATTGTCAGCTCCCCTAACCCC	3626
1453	GGGGUAGGGGAGCUGACAAUUCGU	3023	ACGAATTGTCAGCTCCCCTAACCCC	3627
1454	GGGUUAGGGGAGCUGACAAUUCGUG	3024	CACGAATTGTCAGCTCCCCTAACCC	3628
1455	GGUUAGGGGAGCUGACAAUUCGUGG	3025	CCACGAATTGTCAGCTCCCCTAACC	3629
1456	GUUAGGGGAGCUGACAAUUCGUGGG	3026	CCCACGAATTGTCAGCTCCCCTAAC	3630
1457	UUAGGGGAGCUGACAAUUCGUGGGU	3027	ACCCACGAATTGTCAGCTCCCCTAA	3631
1458	UAGGGGAGCUGACAAUUCGUGGGUC	3028	GACCCACGAATTGTCAGCTCCCCTA	3632
1459	AGGGGAGCUGACAAUUCGUGGGUCC	3029	GGACCCACGAATTGTCAGCTCCCCT	3633
1460	GGGGAGCUGACAAUUCGUGGGUCCG	3030	CGGACCCACGAATTGTCAGCTCCCC	3634
1462	GGAGCUGACAAUUCGUGGGUCCGCA	3031	TGCGGACCCACGAATTGTCAGCTCC	3635
1463	GAGCUGACAAUUCGUGGGUCCGCAA	3032	TTGCGGACCCACGAATTGTCAGCTC	3636
1464	AGCUGACAAUUCGUGGGUCCGCAAA	3033	TTTGCAGGACCCACGAATTGTCAGCT	3637
1465	GCUGACAAUUCGUGGGUCCGCAAAA	3034	TTTGCAGGACCCACGAATTGTCAGC	3638
1466	CUGACAAUUCGUGGGUCCGCAAAAU	3035	ATTTTGCAGGACCCACGAATTGTCAG	3639
1467	UGACAAUUCGUGGGUCCGCAAAAUUC	3036	GATTTTGCAGGACCCACGAATTGTCA	3640
1468	GACAAUUCGUGGGUCCGCAAAAUUCU	3037	AGATTTTGCAGGACCCACGAATTGTC	3641
1469	ACAAUUCGUGGGUCCGCAAAAUUCUU	3038	AAGATTTTGCAGGACCCACGAATTGT	3642
1470	CAAUUCGUGGGUCCGCAAAAUUCUUA	3039	TAAGATTTTGCAGGACCCACGAATTG	3643
1471	AAUUCGUGGGUCCGCAAAAUUCUAA	3040	TTAAGATTTTGCAGGACCCACGAATT	3644
1472	AUUCGUGGGUCCGCAAAAUUCUAAAC	3041	GTTAAGATTTTGCAGGACCCACGAAT	3645
1473	UUCGUGGGUCCGCAAAAUUCUAAACU	3042	AGTTAAGATTTTGCAGGACCCACGAA	3646
1474	UCGUGGGUCCGCAAAAUUCUAAACUA	3043	TAGTTAAGATTTTGCAGGACCCACGA	3647
1475	CGUGGGUCCGCAAAAUUCUAAACUAC	3044	GTAGTTAAGATTTTGCAGGACCCACG	3648
1476	GUGGGUCCGCAAAAUUCUAAACUACC	3045	GGTAGTTAAGATTTTGCAGGACCCAC	3649
1477	UGGGUCCGCAAAAUUCUAAACUACCU	3046	AGGTAGTTAAGATTTTGCAGGACCCA	3650
1478	GGGUCCGCAAAAUUCUAAACUACCUA	3047	TAGGTAGTTAAGATTTTGCAGGACCC	3651
1479	GGUCCGCAAAAUUCUAAACUACCUAA	3048	TTAGGTAGTTAAGATTTTGCAGGACC	3652
1480	GUCCGCAAAAUUCUAAACUACCUAAU	3049	ATTAGGTAGTTAAGATTTTGCAGGAC	3653
1481	UCCGCAAAAUUCUAAACUACCUAAUA	3050	TATTAGGTAGTTAAGATTTTGCAGGA	3654

Input Sequence = PLN

Oligo Length = 25

PLN (Homo sapiens phospholamban (PLN) mRNA; 1635 bp)

Table 31
Table 31: Anti-Her2 Ribozyme and Substrate Sequence

RPI Nos	nt. Position	NCH Ribozyme Sequence	Seq ID Nos	NCH Substrate Sequence	Seq ID Nos
17214	175	asgsasuscca CUGAUGagggccguuagggccGaa Igucuc B		GAGCACC A UGGAGCU	
17215	212	gsasgsagsgcg CUGAUGagggccguuagggccGaa Igaggag B		CUCUCC U CGCCUCC	
17216	261	uscsasusguc CUGAUGagggccguuagggccGaa Iugccgg B		CCGGCAC A GACAUGA	
17217	297	cscsasgsug CUGAUGagggccguuagggccGaa Iucucgg B		CCGAGAC C CACCUGG	
17218	381	ascsasgsugc CUGAUGagggccguuagggccGaa Icauugg B		CCAAUGC C AGCCUGU	
17219	619	cscsasusuca CUGAUGagggccguuagggccGaa Iaucucu B		AGAGAUC U UGAAAGG	
17220	665	csgsugsgsucc CUGAUGagggccguuagggccGaa Iguagca B		UGCUAAC A GGACACG	
17221	943	ususgsasagu CUGAUGagggccguuagggccGaa Iaggcag B		CUGCCUC C ACUUCAA	
17222	944	gsusgsaag CUGAUGagggccguuagggccGaa Igaggca B		UGCCUCC A CUUCAAC	
17223	946	usgsusuga CUGAUGagggccguuagggccGaa Iuggagg B		CCUCCAC U UCAACCA	
17224	993	usgsusgua CUGAUGagggccguuagggccGaa Iugacca B		UGGUCAC C UACAACA	
17225	997	uscsusgsugu CUGAUGagggccguuagggccGaa Iuaggug B		CACCUAC A ACACAGA	
17226	1000	gsusgsuscug CUGAUGagggccguuagggccGaa Iuuguag B		CUACAAC A CAGACAC	
17227	1017	usgsusgscau CUGAUGagggccguuagggccGaa Iacucaa B		UUGAGUC C AUGCCCA	
17228	1078	asgsusasagu CUGAUGagggccguuagggccGaa Iuaggga B		UCCCUAC A ACUACCU	
17229	1201	asgsascscau CUGAUGagggccguuagggccGaa Icacacu B		AGUGUGC U AUGGUCU	
17230	1285	csuscsasca CUGAUGagggccguuagggccGaa Iaucuuc B		GAAGAUC U UUGGGAG	
17231	1395	gsusasasacc CUGAUGagggccguuagggccGaa Iugaucu B		AGAUCAC A GGUUACC	
17232	1413	gsccsasugc CUGAUGagggccguuagggccGaa Iagaugu B		ACAUCUC A GCAUGGC	
17233	1450	asgsusugc CUGAUGagggccguuagggccGaa Iagacag B		CGUCUUC C AGAACCU	
17234	1624	asasgsasgcu CUGAUGagggccguuagggccGaa Iucccag B		CUGGGAC C AGCUCUU	
17235	1787	gsccsasucc CUGAUGagggccguuagggccGaa Igccccc B		CGGGGCC A GGAGUGC	
17236	2001	gsccsasugua CUGAUGagggccguuagggccGaa Iagaggu B		ACCUCUC C UACAUGC	
17237	2002	gsccsasugu CUGAUGagggccguuagggccGaa Igagagg B		CCUCUCC U ACAUGCC	
17238	2055	asgsusugau CUGAUGagggccguuagggccGaa Iggcaag B		CUUGCCC C AUCAACU	

Table 31

17239	2056	csasgsusuga	cUGAuGagggccguuagggccGaa	Igggcaa B	UUGCCCC A UCAACUG
17240	2068	csasgsagsu	cUGAuGagggccguuagggccGaa	Igugcag B	CUGACC C ACUCCUG
17241	2069	ascsasgsag	cUGAuGagggccguuagggccGaa	Iggugca B	UGCACC A CUCCUGU
17242	2295	cscsuagsguu	cUGAuGagggccguuagggccGaa	Igcaucg B	CGAUGCC C AACCAGG
17243	2351	asgsasuscca	cUGAuGagggccguuagggccGaa	Icacuuu B	AAGGUGC U UGGAUCU
17244	2373	usgsusasgac	cUGAuGagggccguuagggccGaa	Iugccaa B	UUGGCAC A GUCUACA
17245	2570	csasusasagc	cUGAuGagggccguuagggccGaa	Iugucac B	GUGACAC A GCUUAUG
17246	2665	asuscscscuu	cUGAuGagggccguuagggccGaa	Igcaauc B	GAUUGCC A AGGGGAU
17247	2702	gsuscscscug	cUGAuGagggccguuagggccGaa	Iuacgag B	CUCGUAC A CAGGGAC
17248	2771	cscsgasagcc	cUGAuGagggccguuagggccGaa	Icccga B	UUCGGGC U GGCUCGG
18261	2780	gsuscscsagc	cUGAuGagggccguuagggccGaa	Iccgagc B	GCUCGGC U GCUGGAC
17249	2783	asasusgsucc	cUGAuGagggccguuagggccGaa	Icagccg B	CGGCUGC U GGACAUU
18265	2783	asusgsusucc	cUGAuGagggccguuagggccGaa	Icagcc B	GGCUGC U GGACAU
18267	2783	asasusgsucc	cUGAuGagggccguuagggccGaa	Icagccg B	CGGCUGC U GGACAUU
18269	2783	asusgsusucc	cUGAuGagggccguuagggccGaa	Icagcc B	GGCUGC U GGACAU
18271	2783	asasusgsucc	cUGAuGagggccguuagggccGaa	Icagccg B	CGGCUGC U GGACAUU
18273	2783	asusgsusucc	cUGAuGagggccguuagggccGaa	Icagcc B	GGCUGC U GGACAU
18275	2783	asasusgsucc	cUGAuGagggccguuagggccGaa	Icagccg B	CGGCUGC U GGACAUU
18277	2783	asusgsusucc	cUGAuGagggccguuagggccGaa	Icagcc B	GGCUGC U GGACAU
18262	2788	uscsuscscaa	cUGAuGagggccguuagggccGaa	Iuccagc B	GCUGGAC A UUGACGA
17250	2799	ggsusascuc	cUGAuGagggccguuagggccGaa	Iucucgu B	ACGAGAC A GAGUACC
18263	2935	usasasgsuu	cUGAuGagggccguuagggccGaa	Igccccca B	UGGGGCC A AACCUUA
17251	2939	asuscsgsuua	cUGAuGagggccguuagggccGaa	Iuuuggc B	GCCAAAC C UUACGAU
18266	2939	uscsusgsuua	cUGAuGagggccguuagggccGaa	Iuuuggc B	CCAAAC C UUACGA
18268	2939	asuscsgsuua	cUGAuGagggccguuagggccGaa	Iuuuggc B	GCCAAAC C UUACGAU
18270	2939	uscsusgsuua	cUGAuGagggccguuagggccGaa	Iuuuggc B	CCAAAC C UUACGA
18272	2939	asuscsgsuua	cUGAuGagggccguuagggccGaa	Iuuuggc B	GCCAAAC C UUACGAU
18274	2939	uscsusgsuua	cUGAuGagggccguuagggccGaa	Iuuuggc B	CCAAAC C UUACGA
18276	2939	asuscsgsuua	cUGAuGagggccguuagggccGaa	Iuuuggc B	GCCAAAC C UUACGAU
18278	2939	uscsusgsuua	cUGAuGagggccguuagggccGaa	Iuuuggc B	CCAAAC C UUACGA
18264	2940	csasuscsgua	cUGAuGagggccguuagggccGaa	Iguuugg B	CCAAACC U UACGAUG

Table 31

17252	3096	asgsasasuuu	cUGAuGagggccguuaggccGaa	Iacacca B	UGGUGUC U GAUUUCU
17253	3423	asuaascsauc	cUGAuGagggccguuaggccGaa	Iagccag B	CUGGCUC C GAUGUUAU
17254	3491	asgsgsgscug	cUGAuGagggccguuaggccGaa	Igucaug B	CAUGACC C CAGCCCU
17255	3497	csuagsusaga	cUGAuGagggccguuaggccGaa	Igcuggg B	CCCAGCC C UCUCACAG
17256	3546	asgscscsauc	cUGAuGagggccguuaggccGaa	Iucucag B	CUGAGAC U GAUGGCU
17257	3669	cscsaasgsagu	cUGAuGagggccguuaggccGaa	Icaccag B	CUGGUGC C ACUCUGG
17258	3690	gsgsgsaasag	cUGAuGagggccguuaggccGaa	Iucuugg B	CCAAGAC U CUCUCCC
17259	3744	uscuscsac	cUGAuGagggccguuaggccGaa	Icaccac B	GGGGUGC C GUGGAGA
17260	3817	asasgsgscug	cUGAuGagggccguuaggccGaa	Icugaag B	CUUCAGC C CAGCCUU
17261	3832	usaaasusaga	cUGAuGagggccguuaggccGaa	Iuugucg B	CGACAAC C UCUAUUA
17262	3998	gsaaasgsuac	cUGAuGagggccguuaggccGaa	Iccuucc B	GGAAAGC C UGACUUC
		Hammerhead Ribozyme Sequence			HH Substrate Sequence
17180	191	gscsgsgscac	cUGAuGagggccguuaggccGaa	Aggccc B	GCGGCCU U GUGCCGC
17202	312	gsguugsgcg	cUGAuGagggccguuaggccGaa	Agcaugu B	ACAUGCU C CGCCACC
17182	495	cscuuscscac	cUGAuGagggccguuaggccGaa	Agcuggg B	CCCAGCU C UUUGAGG
17183	498	usgsuuscscu	cUGAuGagggccguuaggccGaa	Aagagcu B	AGCUCUU U GAGGACA
17181	633	gsgsaascaa	cUGAuGagggccguuaggccGaa	Acccuc B	GAGGGGU C UUGAUCC
17276	831	gsgscsasca	cUGAuGagggccguuaggccGaa	Acagugc B	GCACUGUCUGUGCC
17203	995	usgsuugsguug	cUGAuGagggccguuaggccGaa	Aggugac B	GUCACCU A CAACACA
17184	1076	gsuaasgsuug	cUGAuGagggccguuaggccGaa	Agggaca B	UGUCCCU A CAACUAC
17185	1202	csasgsascca	cUGAuGagggccguuaggccGaa	Agcacac B	GUGUGCU A UGGUCUG
17186	1286	gscsuuscscac	cUGAuGagggccguuaggccGaa	Agauuu B	AAGAUU U UGGGAGC
17187	1287	gsgscsuusccc	cUGAuGagggccguuaggccGaa	Aagaucu B	AGAUCUU U GGGAGCC
17188	1317	cscscscsauc	cUGAuGagggccguuaggccGaa	Aagcucu B	AGAGCUU U GAUGGGG
17204	1412	cscsaasugcu	cUGAuGagggccguuaggccGaa	Agaugua B	UACAUCU C AGCAUGG
17189	1448	gsuauuscsgg	cUGAuGagggccguuaggccGaa	Agacgcu B	AGCGUCU U CCAGAAC
17190	1449	gsgsuauuscug	cUGAuGagggccguuaggccGaa	Aagagc B	CGGUCUU C CAGAACC
17191	1632	gsgsuauusccg	cUGAuGagggccguuaggccGaa	Aagagcu B	AGCUCUU U CGGAACC
17192	1761	usgscaasguu	cUGAuGagggccguuaggccGaa	Acacacu B	AGUGUGU C AACUGCA
17193	2000	csaasusguag	cUGAuGagggccguuaggccGaa	Agagguc B	GACCUCU C CUACAUG

Table 31

17194	2003	gsgsgscsaug	cUGAuGagggccguuaggccGaa	Aggagag B		CUCUCCU A CAUGCCC	
17205	2058	usgscsasguu	cUGAuGagggccguuaggccGaa	Auggggc B		GCCCAU C AACUGCA	
17206	2072	csascsasacag	cUGAuGagggccguuaggccGaa	Agugggu B		ACCCACU C CUGUGUG	
17195	2352	csasgsasucc	cUGAuGagggccguuaggccGaa	Agcaccu B		AGGUGCU U GGAUCUG	
17196	2575	usasgsgsgca	cUGAuGagggccguuaggccGaa	Aagcuu B		ACAGCUU A UGCCCCA	
17197	2697	usgsuagbuac	cUGAuGagggccguuaggccGaa	Agccgca B		UGCGGCU C GUACACA	
18257	2776	asgscsasgcc	cUGAuGagggccguuaggccGaa	Agccagc B		GCUGGCU C GGCUGCU	
18258	2790	uscuscsauc	cUGAuGagggccguuaggccGaa	Augucca B		UGGACAU U GACGAGA	
17207	2926	gscscscscaa	cUGAuGagggccguuaggccGaa	Agucauc B		GAUGACU U UUGGGGC	
18259	2928	usgsgscsccc	cUGAuGagggccguuaggccGaa	Aaaguca B		UGACUUU U GGGGCCA	
18260	2942	cscscsasucg	cUGAuGagggccguuaggccGaa	Aagguuu B		AAACCUU A CGAUGGG	
17198	3089	asgsascsacc	cUGAuGagggccguuaggccGaa	Acucccg B		CGGGAGU U GGUGUCU	
17208	3155	usgsgsgsccc	cUGAuGagggccguuaggccGaa	Aguccuc B		GAGGACU U GGGCCCA	
17209	3499	csgscsausua	cUGAuGagggccguuaggccGaa	Agggcug B		CAGCCCU C UACAGCG	
17210	3501	ascscsgscug	cUGAuGagggccguuaggccGaa	Agagggc B		GCCCUU A CAGCGGU	
17211	3714	csusuuusgac	cUGAuGagggccguuaggccGaa	Accccau B		AUGGGGU C GUCAAAG	
17199	3802	gscsaagsgag	cUGAuGagggccguuaggccGaa	Agggugg B		CCACCCU C CUCCUGC	
17200	3825	gsgsuuusguc	cUGAuGagggccguuaggccGaa	Aaggcug B		CAGCCUU C GACAACC	
17201	3838	uscscscsagu	cUGAuGagggccguuaggccGaa	Auagagg B		CCUCUUAU U ACUGGGA	

Lowercase = 2'-O-methyl Modifications

Uppercase: 1. Uppercase U under Ribozyme Sequence = 2'-C-Allyl U

2. All other Uppercases = Ribonucleotides

I = Inosine

B = 3'-3' Inverted abasic deoxyribose

Table 32

Table 32: RNA Cleavage by NCH-XYLO Ribozyme
(Reaction: 50 mM TRIS-Cl pH 7.5, 10 mM Mg²⁺, 37°C, 500 nM FINAL [Rz], Trace Substrate)

DATA SUMMARY

NCH-Xylo Ribozyme (RPI No.)	TARGET TRIPLET 5'-NCX-3'	k _{obs} (min. ⁻¹) guua + A15.1 = xylo
14827	5'-GCA-3'	1.649
14828	5'-ACA-3'	0.293
14829	5'-UCA-3'	0.272
14830	5'-CCA-3'	0.214

Table 33

Table 33: Examples of NCH-Xylo Ribozyme and Substrate Sequences

RPI Nos.	Ribozyme Sequence		Seq. ID. Nos.	Substrate Sequence	Seq. ID. Nos.
14827	5'-ucu cca u	cUGA uGa ggcc guua ggcc Gaa I cuc ccuB-3'		5'-AGGGA GCA AUGGAGA-3'	
14828	5'-ucu cca u	cUGA uGa ggcc guua ggcc Gaa I uuc ccuB-3'		5'-AGGGA ACA AUGGAGA-3'	
14829	5'-ucu cca u	cUGA uGa ggcc guua ggcc Gaa I auc ccuB-3'		5'-AGGGA UCA AUGGAGA-3'	
14830	5'-ucu cca u	cUGA uGa ggcc guua ggcc Gaa I guc ccuB-3'		5'-AGGGA CCA AUGGAGA-3'	

Uppercase = Ribonucleotides
Lowercase = 2'-O-methyl nucleotides
I = Xylo-Inosine
B = 3'-3' inverted abasic
U = 2'-C-allyl-U

Table 34

Table 34: Anti-HER2 NCH Ribozyme and Target Sequences

nt. Position	NCH Substrate Sequence	Seq. ID Nos.	NCH Ribozyme Sequence	Seq. ID Nos.
14	AGGUAAC C CUGGCCC		GGGCCAG CUGAUGAG X CGAA IUUACCU	
15	GGUAACC C UGGCCCC		GGGGCCA CUGAUGAG X CGAA IGUUAAC	
16	GUAACCC U GGCCCCU		AGGGGCC CUGAUGAG X CGAA IGGUUAAC	
20	CCCUGGC C CCUUGG		CCAAAGG CUGAUGAG X CGAA ICCAGGG	
21	CCUGGCC C CUUUGGU		ACCAAAG CUGAUGAG X CGAA IGCCAGG	
22	CUGGCCC C UUUGGUC		GACCAA CUGAUGAG X CGAA IGGCCAG	
23	UGGCCCC U UUGGUCG		CGACCAA CUGAUGAG X CGAA IGGGCCA	
35	UCGGGGC C CCGGCCA		UGCCCGG CUGAUGAG X CGAA ICCCCGA	
36	CGGGGCC C CGGGCAG		CUGCCCG CUGAUGAG X CGAA IGCCCCG	
37	GGGGCCC C GGGCAGC		GCUGCCC CUGAUGAG X CGAA IGGCCCC	
42	CCCGGGC A GCGCGC		GCGCGGC CUGAUGAG X CGAA ICCCGGG	
45	GGGCAGC C GCGCGCC		GGCGCGC CUGAUGAG X CGAA ICUGCCC	
52	CGCGCGC C CCUCCCC		GGGAAGG CUGAUGAG X CGAA ICGCGCG	
53	GCGCGCC C CUUCCCA		UGGAAG CUGAUGAG X CGAA ICGCGC	
54	CGCGCCC C UJCCAC		GUGGGA CUGAUGAG X CGAA IGGCGCG	
55	GCGCCCC U UCCACG		CGUGGGA CUGAUGAG X CGAA IGGCGC	
58	CCCUUC C CACGGG		CCCCGUG CUGAUGAG X CGAA IAAGGGG	
59	CCCUUC C ACGGGC		GCCCCGU CUGAUGAG X CGAA IGAAGGG	
60	CCUUC C CGGGCC		GGCCCCG CUGAUGAG X CGAA IGAAGG	
67	ACGGGGC C CUUACU		AGUAAAG CUGAUGAG X CGAA ICCCCGU	
68	CGGGGCC C UUUACUG		CAGUAAA CUGAUGAG X CGAA IGCCCCG	
69	GGGGCCC U UUACUG		GCAGUAA CUGAUGAG X CGAA IGGCCCC	
74	CCUUUAC U GCGCGC		GCGGCGC CUGAUGAG X CGAA IUAAAGG	
79	ACUGCGC C GCGCGCC		GGCGCGC CUGAUGAG X CGAA ICGCAGU	
86	CGCGCGC C CGGCCCC		GGGGCCG CUGAUGAG X CGAA ICGCGCG	
87	GCGCGCC C GGCCCC		GGGGGCC CUGAUGAG X CGAA ICGCGC	
91	GCCCGGC C CCCACCC		GGGUGGG CUGAUGAG X CGAA ICCGGGC	
92	CCCGGCC C CCACCC		GGGUGG CUGAUGAG X CGAA IGCCGGG	
93	CCGGCCC C CACCCU		AGGGGUG CUGAUGAG X CGAA IGGCCGG	
94	CGGCCCC C ACCCCU		GAGGGGU CUGAUGAG X CGAA IGGGCCG	
95	GGCCCCC A CCCCUCG		CGAGGGG CUGAUGAG X CGAA IGGGCC	
97	CCCCCAC C CCUCGCA		UGCAGG CUGAUGAG X CGAA IUGGGGG	
98	CCCCACC C CUCGAG		CUCGAG CUGAUGAG X CGAA IGUGGGG	
99	CCACCC C UCGCAGC		GCUCGA CUGAUGAG X CGAA IGGUGGG	
100	CCACCC U CGCAGCA		UGCUGC CUGAUGAG X CGAA IGGUGG	
104	CCUUGC A GCACCC		GGGUGC CUGAUGAG X CGAA ICGAGG	
107	UCGAGC A CCCCGC		CGCGGG CUGAUGAG X CGAA ICUGCGA	
109	GCAGCAC C CCGCGC		GGCGCG CUGAUGAG X CGAA IUGCUG	
110	CAGCACC C CGGCCC		GGGCGC CUGAUGAG X CGAA IGUGCUG	
111	AGCACCC C GCGCCC		GGGCGC CUGAUGAG X CGAA IGGUGCU	
116	CCCGCGC C CCGCGC		GGCGCG CUGAUGAG X CGAA ICGCGG	
117	CCCGCGC C CCGCGC		GGCGCG CUGAUGAG X CGAA ICGCGG	

Table 34

118	CGCGCCC C GCGCCCU		AGGGCGC CUGAUGAG X CGAA IGGCGCG	
123	CCCGCGC C CUCCCAG		CUGGGAG CUGAUGAG X CGAA ICGCGGG	
124	CCGCGCC C UCCAGC		GCUGGGA CUGAUGAG X CGAA ICGCGGG	
125	CGCGCCC U CCCAGCC		GGCUGGG CUGAUGAG X CGAA IGGCGCG	
127	CGCCCUC C CAGCCGG		CCGGCUG CUGAUGAG X CGAA IAGGGCG	
128	GCCCUCC C AGCGGG		CCCGGCU CUGAUGAG X CGAA IGAGGGC	
129	CCCUCCC A GCCGGGU		ACCCGGC CUGAUGAG X CGAA IGAGGGG	
132	UCCCAGC C GGUCCA		UGGACCC CUGAUGAG X CGAA ICUGGGA	
138	CCGGGUC C AGCGGA		UCCGGCU CUGAUGAG X CGAA IACCCGG	
139	CGGGUCC A GCCGGAG		CUCCGGC CUGAUGAG X CGAA IGACCCG	
142	GUCCAGC C GGAGCCA		UGGCUCC CUGAUGAG X CGAA ICUGGAC	
148	CCGGAGC C AUGGGGC		GCCCCAU CUGAUGAG X CGAA ICUCCGG	
149	CGGAGCC A UGGGGCC		GGCCCCA CUGAUGAG X CGAA IGCUCCG	
156	AUGGGGC C GGAGCCG		CGGCUC CUGAUGAG X CGAA ICCCAU	
162	CCGGAGC C GCAGUGA		UCACUGC CUGAUGAG X CGAA ICUCCGG	
165	GAGCCGC A GUGAGCA		UGCUCAC CUGAUGAG X CGAA ICGGCU	
172	AGUGAGC A CCAUGGA		UCCAUGG CUGAUGAG X CGAA ICUCACU	
174	UGAGCAC C AUGGAGC		GCUCCAU CUGAUGAG X CGAA IUGCUCA	
175	GAGCACC A UGGAGCU		AGCUCCA CUGAUGAG X CGAA IGUGCUC	
182	AUGGAGC U GGCGGCC		GGCCGCC CUGAUGAG X CGAA ICUCCAU	
189	UGGCGGC C UUGUGCC		GGCACA CUGAUGAG X CGAA ICCGCCA	
190	GGCGGCC U UGUGCCG		CGGCACA CUGAUGAG X CGAA IGCCGCC	
196	CUUGUGC C GCUGGGG		CCCCAGC CUGAUGAG X CGAA ICACAAG	
199	GUGCCGC U GGGGGCU		AGCCCCC CUGAUGAG X CGAA ICGGCAC	
206	UGGGGGC U CCUCCUC		GAGGAGG CUGAUGAG X CGAA ICCCCCA	
208	GGGGCUC C UCCUCGC		GCGAGGA CUGAUGAG X CGAA IAGCCCC	
209	GGGCUC C CCUCGCC		GGCGAGG CUGAUGAG X CGAA IGAGCCC	
211	GTUCCUC C UGCCCCU		AGGGCGA CUGAUGAG X CGAA IAGGAGC	
212	CUCCUCC U CGCCUC		GAGGGCG CUGAUGAG X CGAA IGAGGAG	
216	UCCUCGC C CUCUUGC		GCAAGAG CUGAUGAG X CGAA ICGAGGA	
217	CCUCGCC C UCUUGCC		GGCAAGA CUGAUGAG X CGAA ICGGAGG	
218	CUCGCC C CUUGCCC		GGGCAAG CUGAUGAG X CGAA IGGCGAG	
220	CGCCCUC U UGCCCCC		GGGGGCA CUGAUGAG X CGAA IAGGGCG	
224	CUCUUGC C CCCCGA		UCCGGGG CUGAUGAG X CGAA ICAAGAG	
225	UCUUGCC C CCCGGAG		CUCCGGG CUGAUGAG X CGAA IGCAAGA	
226	CUUGCCC C CCGAGC		GCUCCGG CUGAUGAG X CGAA IGGCAAG	
227	UUGCCCC C CGGAGCC		GGCUCCG CUGAUGAG X CGAA IGGGCAA	
228	UGCCCCC C GGAGCCG		CGGCUC CUGAUGAG X CGAA IGGGGCA	
234	CCGGAGC C GCGAGCA		UGCUCGC CUGAUGAG X CGAA ICUCCGG	
241	CGCGAGC A CCAAGU		ACUUGGG CUGAUGAG X CGAA ICUCGCG	
243	CGAGCAC C CAAGUGU		ACACUUG CUGAUGAG X CGAA IUGCUCG	
244	GAGCACC C AAGUGUG		CACACU CUGAUGAG X CGAA IGUGCUC	
245	AGCACCC A AGUGUGC		GCACACU CUGAUGAG X CGAA IGGUGCU	
253	AGUGUGC A CCGGCAC		GUGCCGG CUGAUGAG X CGAA ICACACU	
255	UGUGCAC C GGCACAG		CUGUGCC CUGAUGAG X CGAA IUGCACA	
259	CACCGGC A CAGACAU		AUGUCUG CUGAUGAG X CGAA ICCGGUG	
261	CCGGCAC A GACAUGA		UCAUGUC CUGAUGAG X CGAA IUGCCGG	

Table 34

265	CACAGAC A UGAAGCU		AGCUUCA CUGAUGAG X CGAA IUCUGUG	
272	AUGAAGC U GCGGCUC		GAGCCGC CUGAUGAG X CGAA ICUUCAU	
278	CUGCGGC U CCCUGCC		GGCAGGG CUGAUGAG X CGAA ICCGCAG	
280	GCGGCUC C CUGCCAG		CUGGCAG CUGAUGAG X CGAA IAGCCGC	
281	CGGCUCC C UGCCAGU		ACUGGCA CUGAUGAG X CGAA IGAGCCG	
282	GGCUCCC U GCCAGUC		GACUGGC CUGAUGAG X CGAA IGGAGCC	
285	UCCUGGC C AGUCCCG		CGGACU CUGAUGAG X CGAA ICAGGGA	
286	CCCUGCC A GUCCCGA		UCGGGAC CUGAUGAG X CGAA IGCAGGG	
290	GCCAGUC C CGAGACC		GGUCUCG CUGAUGAG X CGAA IACUGGC	
291	CCAGUCC C GAGACCC		GGGUCUC CUGAUGAG X CGAA IGACUGG	
297	CCGAGAC C CACCUGG		CCAGGUG CUGAUGAG X CGAA IUCUCGG	
298	CGAGACC C ACCUGGA		UCCAGGU CUGAUGAG X CGAA IGUCUCG	
299	GAGACCC A CCUGGAC		GUCCAGG CUGAUGAG X CGAA IGGUCUC	
301	GACCCAC C UGGACAU		AUGUCCA CUGAUGAG X CGAA IUGGGUC	
302	ACCCACC U GGACAUG		CAUGUCC CUGAUGAG X CGAA IGUGGGU	
307	CCUGGAC A UGCUCCG		CGGAGCA CUGAUGAG X CGAA IUCCAGG	
311	GACAUGC U CCGCCAC		GUGGCGG CUGAUGAG X CGAA ICAUGUC	
313	CAUGCUC C GCCACCU		AGGUGGC CUGAUGAG X CGAA IAGCAUG	
316	GCUCCGC C ACCUCUA		UAGAGGU CUGAUGAG X CGAA ICGGAGC	
317	CUCCGCC A CCUCUAC		GUAGAGG CUGAUGAG X CGAA ICGGAG	
319	CCGCCAC C UCUACCA		UGGUAGA CUGAUGAG X CGAA IUGGCGG	
320	CGCCACC U CUACCAG		CUGGUAG CUGAUGAG X CGAA IGUGGCG	
322	CCACCUC U ACCAGGG		CCCUGGU CUGAUGAG X CGAA IAGGUGG	
325	CCUCUAC C AGGGCUG		CAGCCCU CUGAUGAG X CGAA IUAGAGG	
326	CUCUACC A GGGCUGC		GCAGCCC CUGAUGAG X CGAA IGUAGAG	
331	CCAGGGC U GCCAGGU		ACCUGGC CUGAUGAG X CGAA ICCUGG	
334	GGGCUGC C AGGUGGU		ACCACCU CUGAUGAG X CGAA ICAGCCC	
335	GGCUGCC A GGUGGUG		CACCACC CUGAUGAG X CGAA IGCAGCC	
344	GUGGUGC A GGGAAAC		GUUCCCC CUGAUGAG X CGAA ICACCAC	
352	GGGAAAC C UGGAACU		AGUUCCA CUGAUGAG X CGAA IUUUCCC	
353	GGAAACC U GGAACUC		GAGUCC CUGAUGAG X CGAA IGUUUCC	
359	CUGGAAC U CACCUC		GUAGGUG CUGAUGAG X CGAA IUUCCAG	
361	GGAACUC A CCUACCU		AGGUAGG CUGAUGAG X CGAA IAGUCC	
363	AACUCAC C UACCUGC		GCAGGUA CUGAUGAG X CGAA IUGAGUU	
364	ACUCACC U ACCUGCC		GGCAGGU CUGAUGAG X CGAA IGUGAGU	
367	CACCUAC C UGCCCAC		GUGGGCA CUGAUGAG X CGAA IUAGGUG	
368	ACCUACC U GCCCACC		GGUGGGC CUGAUGAG X CGAA IGUAGGU	
371	UACCUGC C CACCAAU		AUUGGUG CUGAUGAG X CGAA ICAGGUA	
372	ACCUGCC C ACCAAUG		CAUUGGU CUGAUGAG X CGAA IGCAGGU	
373	CCUGCCC A CCAAUGC		GCAUUGG CUGAUGAG X CGAA IGGCAGG	
375	UGCCCAC C AAUGCCA		UGGCAUU CUGAUGAG X CGAA IUGGGCA	
376	GCCCACC A AUGCCAG		CUGGCAU CUGAUGAG X CGAA IGUGGGC	
381	CCAAUGC C AGCCUGU		ACAGGCU CUGAUGAG X CGAA ICAUUGG	
382	CAAUGCC A GCCUGUC		GACAGGC CUGAUGAG X CGAA IGCAUUG	
385	UGCCAGC C UGUCCUU		AAGGACA CUGAUGAG X CGAA ICUGGCA	
386	GCCAGCC U GUCCUUC		GAAGGAC CUGAUGAG X CGAA IGCUGGC	
390	GCCUGUC C UUCCUGC		GCAGGAA CUGAUGAG X CGAA IACAGGC	

Table 34

391	CCUGUCC U UCCUGCA		UGCAGGA CUGAUGAG X CGAA IGACAGG	
394	GUCCUUC C UGCAGGA		UCCUGCA CUGAUGAG X CGAA IAAGGAC	
395	UCCUUC C U GCAGGAU		AUCCUGC CUGAUGAG X CGAA IGAAGGA	
398	UUCUUGC A GGAUAUC		GAUAUCC CUGAUGAG X CGAA ICAGGAA	
406	GGUAUUC C AGGAGGU		ACCUCU CUGAUGAG X CGAA IAUAUCC	
407	GAUAUCC A GGAGGUG		CACCUCC CUGAUGAG X CGAA IGAUAUC	
416	GAGGUGC A GGGCUAC		GUAGCCC CUGAUGAG X CGAA ICACCUC	
421	GCAGGGC U ACGUGCU		AGCACGU CUGAUGAG X CGAA ICCUGC	
428	UACGUGC U CAUCGCU		AGCGAUG CUGAUGAG X CGAA ICACGUA	
430	CGUGCUC A UCGUCA		UGAGCGA CUGAUGAG X CGAA IAGCAGG	
435	UCAUCGC U CACAACC		GGUUGUG CUGAUGAG X CGAA ICGAUGA	
437	AUCGCUC A CAACCAA		UUGGUUG CUGAUGAG X CGAA IAGCGAU	
439	CGCUCAC A ACCAAGU		ACUUGGU CUGAUGAG X CGAA IUAGGCG	
442	UCACAAC C AAGUGAG		CUCACU CUGAUGAG X CGAA IUUGUGA	
443	CACAACC A AGUGAGG		CCUCACU CUGAUGAG X CGAA IGUUGUG	
452	GUGAGGC A GGUCCCA		UGGGACC CUGAUGAG X CGAA ICCUCAC	
457	GCAGGUC C CACUGCA		UGCAGUG CUGAUGAG X CGAA IACCUGC	
458	CAGGUCC C ACUGCAG		CUGCAGU CUGAUGAG X CGAA IGACCUG	
459	AGGUCCC A CUGCAGA		UCUGCAG CUGAUGAG X CGAA IGGACCU	
461	GUCCAC U GCAGAGG		CCUCUGC CUGAUGAG X CGAA IUUGGAC	
464	CCACUGC A GAGGUG		CAGCCUC CUGAUGAG X CGAA ICAGUGG	
470	CAGAGGC U GCGGAUU		AAUCCGC CUGAUGAG X CGAA ICCUCUG	
487	GCGAGGC A CCCAGCU		AGCUGGG CUGAUGAG X CGAA ICCUCGC	
489	GAGGCAC C CAGCUCU		AGAGCUG CUGAUGAG X CGAA IUGCCUC	
490	AGGCACC C AGCUCU		AAGAGCU CUGAUGAG X CGAA IGUGCCU	
491	GGCACCC A GCUCUUU		AAAGAGC CUGAUGAG X CGAA IGGUGCC	
494	ACCCAGC U CUUUGAG		CUCAAAG CUGAUGAG X CGAA ICUGGGU	
496	CCAGCUC U UUGAGGA		UCCUCA CUGAUGAG X CGAA IAGCUGG	
505	UGAGGAC A ACUAUGC		GCAUAGU CUGAUGAG X CGAA IUCCUCA	
508	GGACAAC U AUGCCCU		AGGGCAU CUGAUGAG X CGAA IUUGUCC	
513	ACUAUGC C CUGGCCG		CGGCCAG CUGAUGAG X CGAA ICAUAGU	
514	CUAUGCC C UGGCCGU		ACGGCCA CUGAUGAG X CGAA IGCAUAG	
515	UAUGCCC U GGCCGUG		CACGGCC CUGAUGAG X CGAA IGGCAUA	
519	CCCUGGC C GUGCUAG		CUAGCAC CUGAUGAG X CGAA ICCAGGG	
524	GCCGUGC U AGACAAU		AUUGUCU CUGAUGAG X CGAA ICACGGC	
529	GCUAGAC A AUGGAGA		UCUCCAU CUGAUGAG X CGAA IUCUAGC	
538	UGGAGAC C CGCUGAA		UUCAGCG CUGAUGAG X CGAA IUCUCCA	
539	GGAGACC C GCUGAAC		GUUCAGC CUGAUGAG X CGAA IGUCUCC	
542	GACCCGC U GAACAAU		AUUGUUC CUGAUGAG X CGAA ICGGGUC	
547	GCUGAAC A AUACCAC		GUGGUAU CUGAUGAG X CGAA IUUCAGC	
552	ACAAUAC C ACCCUG		CAGGGGU CUGAUGAG X CGAA IUAUUGU	
553	CAAUACC A CCCUGU		ACAGGGG CUGAUGAG X CGAA IGUAUUG	
555	AUACCAC C CCUGUCA		UGACAGG CUGAUGAG X CGAA IUUGUAU	
556	UACCACC C CUGUCAC		GUGACAG CUGAUGAG X CGAA IGUGGUA	
557	ACCACCC C UGUCACA		UGUGACA CUGAUGAG X CGAA IGGUGGU	
558	CCACCCC U GUCACAG		CUGUGAC CUGAUGAG X CGAA IGGUGG	
562	CCCUGUC A CAGGGGC		GCCCCUG CUGAUGAG X CGAA IACAGGG	

Table 34

564	CUGUCAC A GGGGCCU		AGGCCCC CUGAUGAG X CGAA IUGACAG	
570	CAGGGGC C UCCCCAG		CUGGGGA CUGAUGAG X CGAA ICCCCUG	
571	AGGGGCC U CCCCAGG		CCUGGGG CUGAUGAG X CGAA IGCCCCU	
573	GGCCUC C CCAGGAG		CUCCUGG CUGAUGAG X CGAA IAGGCC	
574	GGCCUCC C CAGGAGG		CCUCCUG CUGAUGAG X CGAA IGAGGCC	
575	GCCUCCC C AGGAGGC		GCCUCCU CUGAUGAG X CGAA IGGAGGC	
576	CCUCCCC A GGAGGCC		GGCCUCC CUGAUGAG X CGAA IGGGAGG	
583	AGGAGGC C UGCGGGA		UCCCCGA CUGAUGAG X CGAA ICCUCCU	
584	GGAGGCC U GCGGAG		CUCCCGC CUGAUGAG X CGAA IGCCUCC	
593	CGGGAGC U GCAGCUU		AAGCUGC CUGAUGAG X CGAA ICUCCCG	
596	GAGCUGC A GCUUCGA		UCGAAGC CUGAUGAG X CGAA ICAGCUC	
599	CUGCAGC U UCGAAGC		GCUUCGA CUGAUGAG X CGAA ICUGCAG	
607	UCGAAGC C UCACAGA		UCUGUGA CUGAUGAG X CGAA ICUUCGA	
608	CGAAGCC U CACAGAG		CUCUGUG CUGAUGAG X CGAA IGCUUCG	
610	AAGCCUC A CAGAGAU		AUCUCUG CUGAUGAG X CGAA IAGGCUU	
612	GCCUCAC A GAGAUCU		AGAUCUC CUGAUGAG X CGAA IUGAGGC	
619	AGAGAU C UGAAAGG		CCUUUCA CUGAUGAG X CGAA IAUUCUC	
634	AGGGGUC U UGAUCCA		UGGAUCA CUGAUGAG X CGAA IACCCCU	
640	CUUGAUC C AGCGGAA		UUCCGCU CUGAUGAG X CGAA IAUCAAG	
641	UUGAUCC A GCGGAAC		GUUCCGC CUGAUGAG X CGAA IGAUCAA	
649	GCGGAAC C CCCAGCU		AGCUGGG CUGAUGAG X CGAA IUUCCGC	
650	CGGAACC C CCAGCUC		GAGCUGG CUGAUGAG X CGAA IGUUCCG	
651	GGAACCC C CAGCUCU		AGAGCUG CUGAUGAG X CGAA IGGUUC	
652	GAACCCC C AGCUCUG		CAGAGCU CUGAUGAG X CGAA IGGGUUC	
653	AACCCCC A GCUCUGC		GCAGAGC CUGAUGAG X CGAA IGGGGUU	
656	CCCCAGC U CUGCUAC		GUAGCAG CUGAUGAG X CGAA ICUGGGG	
658	CCAGCUC U GCUACCA		UGGUAGC CUGAUGAG X CGAA IAGCUGG	
661	GCUCUGC U ACCAGGA		UCCUGGU CUGAUGAG X CGAA ICAGAGC	
664	CUGCUAC C AGGACAC		GUGUCCU CUGAUGAG X CGAA IUAGCAG	
665	UGCUAAC A GGACACG		CGUGUCC CUGAUGAG X CGAA IGUAGCA	
670	CCAGGAC A CGAUUUU		AAAAUCG CUGAUGAG X CGAA IUCCUGG	
688	GAAGGAC A UCUUCCA		UGGAAGA CUGAUGAG X CGAA IUCCUUC	
691	GGACAUC U UCCACAA		UUGUGGA CUGAUGAG X CGAA IAUGUCC	
694	CAUCUUC C ACAAGAA		UUCUUGU CUGAUGAG X CGAA IAAGAUG	
695	AUCUUC C AAGAAC		GUUCUUG CUGAUGAG X CGAA IGAAGAU	
697	CUUCCAC A AGAACAA		UUGUUCU CUGAUGAG X CGAA IUGGAAG	
703	CAAGAAC A ACCAGCU		AGCUGGU CUGAUGAG X CGAA IUUCUUG	
706	GAACAAC C AGCUGGC		GCCAGCU CUGAUGAG X CGAA IUUGUUC	
707	AACAACC A GCUGGCU		AGCCAGC CUGAUGAG X CGAA IGUUGUU	
710	AACCAGC U GGCUCUC		GAGAGCC CUGAUGAG X CGAA ICUGGUU	
714	AGCUGGC U CUCACAC		GUGUGAG CUGAUGAG X CGAA ICCAGCU	
716	CUGGCUC U CACACUG		CAGUGUG CUGAUGAG X CGAA IAGCCAG	
718	GGCUCUC A CACUGAU		AUCAGUG CUGAUGAG X CGAA IAGAGCC	
720	CUCUCAC A CUGAUAG		CUAUCAG CUGAUGAG X CGAA IUGAGAG	
722	CUCACAC U GAUAGAC		GUCUAUC CUGAUGAG X CGAA IUGUGAG	
730	GAUAGAC A CCAACCG		CGGUUGG CUGAUGAG X CGAA IUCUAUC	
732	UAGACAC C AACCGCU		AGCGGUU CUGAUGAG X CGAA IUGUCUA	

Table 34

733	AGACACC A ACCGCUC		GAGCGGU CUGAUGAG X CGAA IGUGUCU	
736	CACCAAC C GCUCUCG		CGAGAGC CUGAUGAG X CGAA IUUGGUG	
739	CAACCGC U CUCGGGC		GCCCGAG CUGAUGAG X CGAA ICGGUUG	
741	ACCGCUC U CGGGCCU		AGGCCCG CUGAUGAG X CGAA IAGCGGU	
747	CUCGGGC C UGCCACC		GGUGGCA CUGAUGAG X CGAA ICCCGAG	
748	UCGGGCC U GCCACCC		GGUGGC CUGAUGAG X CGAA IGCCCGA	
751	GGCCUGC C ACCCCUG		CAGGGGU CUGAUGAG X CGAA ICAGGCC	
752	GCCUGCC A CCCUGU		ACAGGGG CUGAUGAG X CGAA IGCAGGC	
754	CUGCCAC C CCUGUUC		GAACAGG CUGAUGAG X CGAA IUGGCAG	
755	UGCCACC C CUGUUCU		AGAACAG CUGAUGAG X CGAA IGUGGCA	
756	GCCACCC C UGUUCUC		GAGAAC A CUGAUGAG X CGAA IGGUGGC	
757	CCACCCC U GUUCUCC		GGAGAAC CUGAUGAG X CGAA IGGUGGG	
762	CCUGUUC U CCGAUGU		ACAUCGG CUGAUGAG X CGAA IAAACAGG	
764	UGUUCUC C GAUGUGU		ACACAUC CUGAUGAG X CGAA IAGAACA	
778	UAAGGGC U CCCGCUG		CAGCGGG CUGAUGAG X CGAA ICCCUUA	
780	AGGGCUC C CGCUGCU		AGCAGCG CUGAUGAG X CGAA IAGCCCU	
781	GGGCUC C GCUCUG		CAGCAGC CUGAUGAG X CGAA IGAGCCC	
784	CUCCCGC U GCUGGGG		CCCCAGC CUGAUGAG X CGAA ICGGGAG	
787	CCGCUGC U GGGGAGA		UCUCCCC CUGAUGAG X CGAA ICAGCGG	
801	AGAGUUC U GAGGAUU		AAUCCUC CUGAUGAG X CGAA IAAUCUCU	
812	GAUUGUC A GAGCCUG		CAGGCUC CUGAUGAG X CGAA IACAAUC	
817	UCAGAGC C UGACGCG		CGCGUCA CUGAUGAG X CGAA ICUCUGA	
818	CAGAGCC U GACGCGC		GCGCGUC CUGAUGAG X CGAA IGCUCUG	
826	GACGCGC A CUGUCUG		CAGACAG CUGAUGAG X CGAA ICGCGUC	
828	CGCGCAC U GUCUGUG		CACAGAC CUGAUGAG X CGAA IUCCGCG	
832	CACUGUC U GUGCCGG		CCGGCAC CUGAUGAG X CGAA IACAGUG	
837	UCUGUGC C GGUGGCU		AGCCACC CUGAUGAG X CGAA ICACAGA	
844	CGGUGGC U GUGCCCG		CGGGCAC CUGAUGAG X CGAA ICCACCG	
849	GCUGUGC C CGCUGCA		UGCAGCG CUGAUGAG X CGAA ICACAGC	
850	CUGUGCC C GCUGCAA		UUGCAGC CUGAUGAG X CGAA IGCACAG	
853	UGCCCGC U GCAAGGG		CCCUUGC CUGAUGAG X CGAA ICGGGCA	
856	CCGCUGC A AGGGGCC		GGCCCCU CUGAUGAG X CGAA ICAGCGG	
863	AAGGGGC C ACUGCCC		GGGCAGU CUGAUGAG X CGAA ICCCCUU	
864	AGGGGCC A CUGCCCA		UGGGCAG CUGAUGAG X CGAA IGCCCUU	
866	GGGCCAC U GCCCACU		AGUGGGC CUGAUGAG X CGAA IUGGCCC	
869	CCACUGC C CACUGAC		GUCAGUG CUGAUGAG X CGAA ICAGUGG	
870	CACUGCC C ACUGACU		AGUCAGU CUGAUGAG X CGAA IGCAGUG	
871	ACUGCCC A CUGACUG		CAGUCAG CUGAUGAG X CGAA IGGCAGU	
873	UGCCAC U GACUGCU		AGCAGUC CUGAUGAG X CGAA IUGGGCA	
877	CACUGAC U GCUGCCA		UGGCAGC CUGAUGAG X CGAA IUCAGUG	
880	UGACUGC U GCCAUGA		UCAUGGC CUGAUGAG X CGAA ICAGUCA	
883	CUGCUGC C AUGAGCA		UGCUCAU CUGAUGAG X CGAA ICAGCAG	
884	UGCUGCC A UGAGCAG		CUGCUC A CUGAUGAG X CGAA IGCAGCA	
890	CAUGAGC A GUGUGCU		AGCACAC CUGAUGAG X CGAA ICUCAUG	
897	AGUGUGC U GCCGGCU		AGCCGGC CUGAUGAG X CGAA ICACACU	
900	GUGCUGC C GGCUGCA		UGCAGCC CUGAUGAG X CGAA ICAGCAC	
904	UGCCGGC U GCACGGG		CCCUGUC CUGAUGAG X CGAA ICCGGCA	

Table 34

907	CGGCUGC A CGGGCCC	GGGCCCC CUGAUGAG X CGAA ICAGCCG
913	CACGGGC C CCAAGCA	UGCUGG CUGAUGAG X CGAA ICCCGUG
914	ACGGGCC C CAAGCAC	GUGCUUG CUGAUGAG X CGAA IGCCCGU
915	CGGGCCC C AAGCACU	AGUGCUU CUGAUGAG X CGAA IGGCCCC
916	GGGCCCC A AGCACUC	GAGUGCU CUGAUGAG X CGAA IGGGCC
920	CCCAAGC A CUCUGAC	GUCAGAG CUGAUGAG X CGAA ICUUGGG
922	CAAGCAC U CUGACUG	CAGUCAG CUGAUGAG X CGAA IUGCUUG
924	AGCACUC U GACUGCC	GGCAGUC CUGAUGAG X CGAA IAGUGCU
928	CUCUGAC U GCCUGGC	GCCAGGC CUGAUGAG X CGAA IUCAGAG
931	UGACUGC C UGGCCUG	CAGGCCA CUGAUGAG X CGAA ICAGUCA
932	GACUGCC U GGCCUGC	GCAGGCC CUGAUGAG X CGAA IGCAGUC
936	GCCUGGC C UGCCUCC	GGAGGCA CUGAUGAG X CGAA ICCAGGC
937	CCUGGCC U GCCUCCA	UGGAGGC CUGAUGAG X CGAA IGCCAGG
940	GGCCUGC C UCCACUU	AAGUGGA CUGAUGAG X CGAA ICAGGCC
941	GCCUGCC U CCACUUC	GAAGUGG CUGAUGAG X CGAA IGCAGGC
943	CUGCCUC C ACUUCAA	UUGAAGU CUGAUGAG X CGAA IAGGCAG
944	UGCCUCC A CUUCAAC	GUUGAAG CUGAUGAG X CGAA IGAGGCA
946	CCUCCAC U UCAACCA	UGGUUGA CUGAUGAG X CGAA IUGGAGG
949	CCACUUC A ACCACAG	CUGUGGU CUGAUGAG X CGAA IAAGUGG
952	CUUCAAC C ACAGUGG	CCACUGU CUGAUGAG X CGAA IUUGAAG
953	UUCAACC A CAGUGGC	GCCACUG CUGAUGAG X CGAA IGUGGAA
955	CAACCAC A GUGGCAU	AUGCCAC CUGAUGAG X CGAA IUGGUUG
961	CAGUGGC A UCUGUGA	UCACAGA CUGAUGAG X CGAA ICCACUG
964	UGGCAUC U GUGAGCU	AGCUCAC CUGAUGAG X CGAA IAUGCCA
971	UGUGAGC U GCACUGC	GCAGUGC CUGAUGAG X CGAA ICUCACA
974	GAGCUGC A CUGCCCA	UGGGCAG CUGAUGAG X CGAA ICAGCUC
976	GCUGCAC U GCCCAGC	GCUGGGC CUGAUGAG X CGAA IUGCAGC
979	GCACUGC C CAGCCCU	AGGGCUG CUGAUGAG X CGAA ICAGUGC
980	CACUGCC C AGCCUG	CAGGGCU CUGAUGAG X CGAA IGCAUG
981	ACUGCCC A GCCCUGG	CCAGGGC CUGAUGAG X CGAA IGGCAGU
984	GCCCAGC C CUGGUCA	UGACCAG CUGAUGAG X CGAA ICUGGGC
985	CCCAGCC C UGGUCAC	GUGACCA CUGAUGAG X CGAA IGCUGGG
986	CCAGCCC U GGUCACC	GGUGACC CUGAUGAG X CGAA IGGCUGG
991	CCUGGUC A CCUACAA	UUGUAGG CUGAUGAG X CGAA IACCAGG
993	UGGUCAC C UACAACA	UGUUGUA CUGAUGAG X CGAA IUGACCA
994	GGUCACC U ACAACAC	GUGUUGU CUGAUGAG X CGAA IGUGACC
997	CACCUAC A ACACAGA	UCUGUGU CUGAUGAG X CGAA IUAGGUG
1000	CUACAAC A CAGACAC	GUGUCUG CUGAUGAG X CGAA IUUGUAG
1002	ACAACAC A GACAGU	ACGUGUC CUGAUGAG X CGAA IUGUUGU
1006	CACAGAC A CGUUGA	UCAACG CUGAUGAG X CGAA IUCUGUG
1017	UUGAGUC C AUGCCCA	UGGGCAU CUGAUGAG X CGAA IACUCAA
1018	UGAGUCC A UGCCCAA	UUGGGCA CUGAUGAG X CGAA IGACUCA
1022	UCCAUGC C CAAUCCC	GGGAUUG CUGAUGAG X CGAA ICAUGGA
1023	CCAUGCC C AAUCCCG	CGGGAUU CUGAUGAG X CGAA IGCAUGG
1024	CAUGCCC A AUCCCGA	UCGGGAU CUGAUGAG X CGAA IGGCAUG
1028	CCCAUUC C CGAGGGC	GCCCUCG CUGAUGAG X CGAA IAUUGGG
1029	CCAAUCC C GAGGGCC	GGCCCUC CUGAUGAG X CGAA IGAUUGG

Table 34

1036	CGAGGGC C GGUAUAC		GUAUACC CUGAUGAG X CGAA ICCUCG	
1044	GGUAUAC A UUCGGCG		CGCCGAA CUGAUGAG X CGAA IUUAUACC	
1053	UCGGCGC C AGCUGUG		CACAGCU CUGAUGAG X CGAA ICGCCGA	
1054	CGGCGCC A GCUGUGU		ACACAGC CUGAUGAG X CGAA ICGCCCG	
1057	CGCCAGC U GUGUGAC		GUCACAC CUGAUGAG X CGAA ICUGGCG	
1065	GUGUGAC U GCCUGUC		GACAGGC CUGAUGAG X CGAA IUCACAC	
1068	UGACUGC C UGUCCCU		AGGGACA CUGAUGAG X CGAA ICAGUCA	
1069	GACUGCC U GUCCCUA		UAGGGAC CUGAUGAG X CGAA IGCAGUC	
1073	GCCUGUC C CUACAAC		GUUGUAG CUGAUGAG X CGAA IACAGGC	
1074	CCUGUCC C UACAACU		AGUUGUA CUGAUGAG X CGAA IGACAGG	
1075	CUGUCCC U ACAACUA		UAGUUGU CUGAUGAG X CGAA IGGACAG	
1078	UCCCUAC A ACUACCU		AGGUAGU CUGAUGAG X CGAA IUAGGGA	
1081	CUACAAC U ACCUUUC		GAAAGGU CUGAUGAG X CGAA IUUGUAG	
1084	CAACUAC C UUCUAC		GUAGAAA CUGAUGAG X CGAA IUAGUUG	
1085	AACUACC U UUCUACG		CGUAGAA CUGAUGAG X CGAA IGUAGUU	
1089	ACCUUUC U ACGGACG		CGUCCGU CUGAUGAG X CGAA IAAAGGU	
1104	UGGGAUC C UGCACCC		GGGUGCA CUGAUGAG X CGAA IAUCCCA	
1105	GGGAUCC U GCACCCU		AGGGUGC CUGAUGAG X CGAA IGAUCCC	
1108	AUCCUGC A CCCUCGU		ACGAGGG CUGAUGAG X CGAA ICAGGAU	
1110	CCUGCAC C CUCGUCU		AGACGAG CUGAUGAG X CGAA IUGCAGG	
1111	CUGCACC C UCGUCUG		CAGACGA CUGAUGAG X CGAA IGUCGAG	
1112	UGCACCC U CGUCUGC		GCAGACG CUGAUGAG X CGAA IGGUGCA	
1117	CCUCGUC U GCCCCCU		AGGGGGC CUGAUGAG X CGAA IACGAGG	
1120	CGUCUGC C CCCUGCA		UGCAGGG CUGAUGAG X CGAA ICAGACG	
1121	GUCUGCC C CCUGCAC		GUGCAGG CUGAUGAG X CGAA IGACAGC	
1122	UCUGCCC C CUGCACA		UGUGCAG CUGAUGAG X CGAA IGGCAGA	
1123	CUGCCCC C UGCACAA		UUGUGCA CUGAUGAG X CGAA IGGGCAG	
1124	UGCCCCC U GCACAAC		GUUGUGC CUGAUGAG X CGAA IGGGGCA	
1127	CCCCUGC A CAACCAA		UUGGUUG CUGAUGAG X CGAA ICAGGGG	
1129	CCUGCAC A ACCAAGA		UCUUGGU CUGAUGAG X CGAA IUGCAGG	
1132	GCACAAC C AAGAGGU		ACCUCUU CUGAUGAG X CGAA IUUGUGC	
1133	CACAACC A AGAGGUG		CACCUCU CUGAUGAG X CGAA IGUUGUG	
1143	AGGUGAC A GCAGAGG		CCUCUGC CUGAUGAG X CGAA IUCACCU	
1146	UGACAGC A GAGGAUG		CAUCCUC CUGAUGAG X CGAA ICUGUCA	
1158	AUGGAAC A CAGCGGU		ACCGCUG CUGAUGAG X CGAA IUUCCAU	
1160	GGAACAC A GCGGUGU		ACACCGC CUGAUGAG X CGAA IUUGUCC	
1177	GAAGUGC A GCAAGCC		GGCUUGC CUGAUGAG X CGAA ICACUUC	
1180	GUGCAGC A AGCCUG		CAGGGCU CUGAUGAG X CGAA ICUGCAC	
1184	AGCAAGC C CUGUGCC		GGCACAG CUGAUGAG X CGAA ICUUGCU	
1185	GCAAGCC C UGUGCCC		GGGCACA CUGAUGAG X CGAA IGCUUGC	
1186	CAAGCCC U GUGCCCG		CGGGCAC CUGAUGAG X CGAA IGGCUUG	
1191	CCUGUGC C CGAGUGU		ACACUCG CUGAUGAG X CGAA ICACAGG	
1192	CUGUGCC C GAGUGUG		CACACUC CUGAUGAG X CGAA IGCACAG	
1201	AGUGUGC U AUGGUCU		AGACCAU CUGAUGAG X CGAA ICACACU	
1208	UAUGGUC U GGGCAUG		CAUGCCC CUGAUGAG X CGAA IACCAUA	
1213	UCUGGGC A UGGAGCA		UGCUGCA CUGAUGAG X CGAA ICCAGA	
1220	AUGGAGC A CUUGCGA		UCGCAAG CUGAUGAG X CGAA ICUCCAU	

Table 34

1222	GGAGCAC U UGCGAGA		UCUCGCA CUGAUGAG X CGAA IUGCUCC	
1239	UGAGGGC A GUUACCA		UGGUAAC CUGAUGAG X CGAA ICCCUCA	
1245	CAGUUAC C AGUGCCA		UGGCACU CUGAUGAG X CGAA IUAACUG	
1246	AGUUACC A GUGCCAA		UUGGCAC CUGAUGAG X CGAA IGUAACU	
1251	CCAGUGC C AAUAUCC		GGUAUUA CUGAUGAG X CGAA ICACUGG	
1252	CAGUGCC A AUAUCCA		UGGAUUA CUGAUGAG X CGAA IGCACUG	
1258	CAUAUAC C AGGAGUU		AACUCCU CUGAUGAG X CGAA IAUUUUG	
1259	AAUAUCC A GGAGUUU		AAACUCC CUGAUGAG X CGAA IGAUUAU	
1269	AGUUUGC U GGCUGCA		UGCAGCC CUGAUGAG X CGAA ICAAACU	
1273	UGCUGGC U GCAAGAA		UUCUUGC CUGAUGAG X CGAA ICCAGCA	
1276	UGGCGUC A AGAAGAU		AUCUUCU CUGAUGAG X CGAA ICAGCCA	
1285	GAAGAUC U UUGGGAG		CUCCCAA CUGAUGAG X CGAA IAUUUUC	
1294	UGGGAGC C UGGCAUU		AAUGCCA CUGAUGAG X CGAA ICUCCCA	
1295	GGGAGCC U GGCAUUU		AAAUGCC CUGAUGAG X CGAA IGCUCCC	
1299	GCCUGGC A UUUCUGC		GCAGAAA CUGAUGAG X CGAA ICCAGGC	
1304	GCAUUUC U GCCGGAG		CUCCGGC CUGAUGAG X CGAA IAAUUGC	
1307	UUUCUGC C GGAGAGC		GCUCUCC CUGAUGAG X CGAA ICAGAAA	
1315	GGAGAGC U UUGAUGG		CCAACAA CUGAUGAG X CGAA ICUCUCC	
1327	UGGGGAC C CAGCCUC		GAGGCUG CUGAUGAG X CGAA IUCCCCA	
1328	GGGGACC C AGCCUCC		GGAGGCU CUGAUGAG X CGAA IGUCCCC	
1329	GGGACCC A GCCUCCA		UGGAGGC CUGAUGAG X CGAA IGGUCCC	
1332	ACCCAGC C UCCAACA		UGUUGGA CUGAUGAG X CGAA ICUGGGU	
1333	CCCAGCC U CCAACAC		GUGUUGG CUGAUGAG X CGAA IGCUGGG	
1335	CAGCCUC C AACACUG		CAGUGUU CUGAUGAG X CGAA IAGGCUG	
1336	AGCCUCC A ACACUGC		GCAGUGU CUGAUGAG X CGAA IGAGGCU	
1339	CUCCAAC A CUGCCCC		GGGGCAG CUGAUGAG X CGAA IUUGGAG	
1341	CCAACAC U GCCCCGC		GCGGGGC CUGAUGAG X CGAA IUGUUGG	
1344	ACACUGC C CCGCUCC		GGAGCGG CUGAUGAG X CGAA ICAGUGU	
1345	CACUGCC C CGCUCCA		UGGAGCG CUGAUGAG X CGAA IGCAGUG	
1346	ACUGCCC C GCUCCAG		CUGGAGC CUGAUGAG X CGAA IGGCAGU	
1349	GCCCCGC U CCAGCCA		UGGCUGG CUGAUGAG X CGAA ICGGGGC	
1351	CCCGCUC C AGCCAGA		UCUGGCU CUGAUGAG X CGAA IAGCGGG	
1352	CCGCUCC A GCCAGAG		CUCUGGC CUGAUGAG X CGAA IGAGCGG	
1355	CUCCAGC C AGAGCAG		CUGCUCU CUGAUGAG X CGAA ICUGGAG	
1356	UCCAGCC A GAGCAGC		GCUGCUC CUGAUGAG X CGAA IGCUGGA	
1361	CCAGAGC A GCUCCAA		UUGGAGC CUGAUGAG X CGAA ICUCUGG	
1364	GAGCAGC U CCAAGUG		CACUUGG CUGAUGAG X CGAA ICUGCUC	
1366	GCAGCUC C AAGUGUU		AACACUU CUGAUGAG X CGAA IAGCUGC	
1367	CAGCUC C AAGUGUU		AAACACU CUGAUGAG X CGAA IGAGCUG	
1380	UUGAGAC U CUGGAAG		CUUCCAG CUGAUGAG X CGAA IUCUCAA	
1382	GAGACUC U GGAAGAG		CUCUUC CUGAUGAG X CGAA IAGUCUC	
1393	AGAGAUC A CAGGUUA		UAACCUG CUGAUGAG X CGAA IAUUCUC	
1395	AGAUCAC A GGUUACC		GGUAACC CUGAUGAG X CGAA IUGAUCU	
1402	AGGUUAC C UAUACAU		AUGUAUA CUGAUGAG X CGAA IUAACCU	
1403	GGUUACC U AUACAUC		GAUGUAU CUGAUGAG X CGAA IGUAACC	
1408	CCUAUAC A UCUCAGC		GCUGAGA CUGAUGAG X CGAA IUAUAGG	
1411	AUACAUC U CAGCAUG		CAUGCUG CUGAUGAG X CGAA IAUGUAU	

Table 34

1413	ACAUCUC A GCAUGGC		GCCAUGC CUGAUGAG X CGAA IAGAUGU	
1416	UCUCAGC A UGGCCGG		CCGGCCA CUGAUGAG X CGAA ICUGAGA	
1421	GCAUGGC C GGACAGC		GCUGUCC CUGAUGAG X CGAA ICCAUGC	
1426	GCCGGAC A GCCUGCC		GGCAGGC CUGAUGAG X CGAA IUCCGGC	
1429	GGACAGC C UGCCUGA		UCAGGCA CUGAUGAG X CGAA ICUGUCC	
1430	GACAGCC U GCCUGAC		GUCAGGC CUGAUGAG X CGAA IGCUGUC	
1433	AGCCUGC C UGACCUC		GAGGUCA CUGAUGAG X CGAA ICAGGCU	
1434	GCCUGCC U GACCUCA		UGAGGUC CUGAUGAG X CGAA IGCAGGC	
1438	GCCUGAC C UCAGCGU		ACGCUGA CUGAUGAG X CGAA IUCAGGC	
1439	CCUGACC U CAGCGUC		GACGCUG CUGAUGAG X CGAA IGUCAGG	
1441	UGACCUC A GCGUCUU		AAGACGC CUGAUGAG X CGAA IAGGUCA	
1447	CAGCGUC U UCCAGAA		UUCUGGA CUGAUGAG X CGAA IACGCUG	
1450	CGUCUUC C AGAACCU		AGGUUCU CUGAUGAG X CGAA IAAGACG	
1451	GUCUUC C AGAACCU		CAGGUUC CUGAUGAG X CGAA IGAAGAC	
1456	CCAGAAC C UGCAAGU		ACUUGCA CUGAUGAG X CGAA IUUCUGG	
1457	CAGAACC U GCAAGUA		UACUUGC CUGAUGAG X CGAA IGUUCUG	
1460	AACCUGC A AGUAAUC		GAUUACU CUGAUGAG X CGAA ICAGGUU	
1468	AGUAAUC C GGGGACG		CGUCCCC CUGAUGAG X CGAA IAUUACU	
1481	CGAAUUC U GCACAAU		AUUGUGC CUGAUGAG X CGAA IAAUUCG	
1484	AUUCUGC A CAAUGGC		GCCAUUG CUGAUGAG X CGAA ICAGAAU	
1486	UCUGCAC A AUGGCGC		GCGCCAU CUGAUGAG X CGAA IUGCAGA	
1494	AUGGCGC C UACUCGC		GCGAGUA CUGAUGAG X CGAA ICGCCAU	
1495	UGGCGCC U ACUCGCU		AGCGAGU CUGAUGAG X CGAA ICGCCA	
1498	CGCCUAC U CGCUGAC		GUCAGCG CUGAUGAG X CGAA IUAGGCG	
1502	UACUCGC U GACCCUG		CAGGGUC CUGAUGAG X CGAA ICGAGUA	
1506	CGCUGAC C CUGCAAG		CUUGCAG CUGAUGAG X CGAA IUCAGCG	
1507	GCUGACC C UGCAAGG		CCUUGCA CUGAUGAG X CGAA IGUCAGC	
1508	CUGACCC U GCAAGGG		CCCUUGC CUGAUGAG X CGAA IGGUCAG	
1511	ACCCUGC A AGGGCUG		CAGCCCU CUGAUGAG X CGAA ICAGGGU	
1517	CAAGGGC U GGGCAUC		GAUGCCC CUGAUGAG X CGAA ICCCUUG	
1522	GCUGGGC A UCAGCUG		CAGCUGA CUGAUGAG X CGAA ICCCAGC	
1525	GGGCAUC A GCUGGCU		AGCCAGC CUGAUGAG X CGAA IAUGCCC	
1528	CAUCAGC U GGCUGGG		CCCAGCC CUGAUGAG X CGAA ICUGAUG	
1532	AGCUGGC U GGGGUG		CAGCCCC CUGAUGAG X CGAA ICCAGCU	
1538	CUGGGGC U GCGCUCA		UGAGCGC CUGAUGAG X CGAA ICCCCAG	
1543	GCUGCGC U CACUGAG		CUCAGUG CUGAUGAG X CGAA ICGCAGC	
1545	UGCGCUC A CUGAGGG		CCCUAG CUGAUGAG X CGAA IAGCGCA	
1547	CGCUCAC U GAGGGAA		UUCUUC CUGAUGAG X CGAA IUGAGCG	
1556	AGGGAAC U GGGCAGU		ACUGCCC CUGAUGAG X CGAA IUUCCCU	
1561	ACUGGGC A GUGGACU		AGUCCAC CUGAUGAG X CGAA ICCCAGU	
1568	AGUGGAC U GGCCUC		GAGGGCC CUGAUGAG X CGAA IUCCACU	
1572	GACUGGC C CUCAUCC		GGAUGAG CUGAUGAG X CGAA ICCAGUC	
1573	ACUGGCC C UCAUCCA		UGGAUGA CUGAUGAG X CGAA IGCCAGU	
1574	CUGGGCC U CAUCCAC		GUGGAUG CUGAUGAG X CGAA IGGCCAG	
1576	GGCCUC A UCCACCA		UGGUGGA CUGAUGAG X CGAA IAGGGCC	
1579	CCUCAUC C ACCAUAA		UUAUGGU CUGAUGAG X CGAA IAUGAGG	
1580	CUCAUCC A CCAUAAC		GUUAUGG CUGAUGAG X CGAA IGAUGAG	

Table 34

1582	CAUCCAC C AUAACAC		GUGUUAU CUGAUGAG X CGAA IUGGAUG	
1583	AUCCACC A UAACACC		GGUGUUA CUGAUGAG X CGAA IGUGGAU	
1588	CCAUAAAC A CCCACCU		AGGUGGG CUGAUGAG X CGAA IUUAUGG	
1590	AUAACAC C CACCUCU		AGAGGUG CUGAUGAG X CGAA IUGUUAU	
1591	UAACACC C ACCUCUG		CAGAGGU CUGAUGAG X CGAA IGUGUUA	
1592	AACACCC A CCUCUGC		GCAGAGG CUGAUGAG X CGAA IGGUGUU	
1594	CACCCAC C UCUGCUU		AAGCAGA CUGAUGAG X CGAA IUGGGUG	
1595	ACCCACC U CUGCUUC		GAAGCAG CUGAUGAG X CGAA IGUGGGU	
1597	CCACCUC U GCUUCGU		ACGAAGC CUGAUGAG X CGAA IAGGUGG	
1600	CCUCUGC U UCGUGCA		UGCACGA CUGAUGAG X CGAA ICAGAGG	
1607	UUCGUGC A CACGGUG		CACCGUG CUGAUGAG X CGAA ICACGAA	
1609	CGUGCAC A CGGUGCC		GGCACCG CUGAUGAG X CGAA IUGCACG	
1616	ACGGUGC C CUGGGAC		GUCCCAG CUGAUGAG X CGAA ICACCGU	
1617	CGGUGCC C UGGGACC		GGUCCCA CUGAUGAG X CGAA IGCACCG	
1618	GGUGCCC U GGGACCA		UGGUCCC CUGAUGAG X CGAA IGGCACC	
1624	CUGGGAC C AGCUCUU		AAGAGCU CUGAUGAG X CGAA IUCCCAG	
1625	UGGGACC A GCUCUUU		AAAGAGC CUGAUGAG X CGAA IGUCCCA	
1628	GACCAGC U CUUUCGG		CCGAAAG CUGAUGAG X CGAA ICUGGUC	
1630	CCAGCUC U UUCGGAA		UUCCGAA CUGAUGAG X CGAA IAGCUGG	
1639	UCGGAAC C CGCACCA		UGGUGCG CUGAUGAG X CGAA IUUCCGA	
1640	CGGAACC C GCACCAA		UUGGUGC CUGAUGAG X CGAA IGUCCG	
1643	AACCCGC A CCAAGCU		AGCUUGG CUGAUGAG X CGAA ICGGGUU	
1645	CCCGCAC C AAGCUCU		AGAGCUU CUGAUGAG X CGAA IUGCGGG	
1646	CCGCACC A AGCUCUG		CAGAGCU CUGAUGAG X CGAA IGUGCGG	
1650	ACCAAGC U CUGCUCC		GGAGCAG CUGAUGAG X CGAA ICUUGGU	
1652	CAAGCUC U GCUCCAC		GUGGAGC CUGAUGAG X CGAA IAGCUUG	
1655	GCUCUGC U CCACACU		AGUGUGG CUGAUGAG X CGAA ICAGAGC	
1657	UCUGCUC C ACACUGC		GCAGUGU CUGAUGAG X CGAA IAGCAGA	
1658	CUGCUCC A CACUGCC		GGCAGUG CUGAUGAG X CGAA IGAGCAG	
1660	GCUCCAC A CUGCCAA		UUGGCAG CUGAUGAG X CGAA IUGGAGC	
1662	UCCACAC U GCCAACC		GGUUGGC CUGAUGAG X CGAA IUGUGGA	
1665	ACACUGC C AACCGGC		GCCGGUU CUGAUGAG X CGAA ICAGUGU	
1666	CACUGCC A ACCGGCC		GGCCGGU CUGAUGAG X CGAA IGCAGUG	
1669	UGCCAAC C GGCCAGA		UCUGGCC CUGAUGAG X CGAA IUUGGCA	
1673	AACCGGC C AGAGGAC		GUCCUCU CUGAUGAG X CGAA ICCGGUU	
1674	ACCGGCC A GAGGACG		CGUCCUC CUGAUGAG X CGAA IGCCGGU	
1699	CGAGGGC C UGGCCUG		CAGGCCA CUGAUGAG X CGAA ICCCUCG	
1700	GAGGGCC U GGCCUGC		GCAGGCC CUGAUGAG X CGAA IGCCCUC	
1704	GCCUGGC C UGCCACC		GGUGGCA CUGAUGAG X CGAA ICCAGGC	
1705	CCUGGCC U GCCACCA		UGGUGGC CUGAUGAG X CGAA IGCCAGG	
1708	GGCCUGC C ACCAGCU		AGCUGGU CUGAUGAG X CGAA ICAGGCC	
1709	GCCUGCC A CCAGCUG		CAGCUGG CUGAUGAG X CGAA IGCAGGC	
1711	CUGCCAC C AGCUGUG		CACAGCU CUGAUGAG X CGAA IUGGCAG	
1712	UGCCACC A GCUGUGC		GCACAGC CUGAUGAG X CGAA IGUGGCA	
1715	CACCAGC U GUGCGCC		GGCGCAC CUGAUGAG X CGAA ICUGGUG	
1722	UGUGCGC C CGAGGGC		GCCUCUG CUGAUGAG X CGAA ICGACA	
1723	GUGCGEC C GAGGGCA		UGCCCUC CUGAUGAG X CGAA IGCACAC	

Table 34

1730	CGAGGGC A CUGCUGG		CCAGCAG CUGAUGAG X CGAA ICCCUUG	
1732	AGGGCAC U GCUGGGG		CCCCAGC CUGAUGAG X CGAA IUGCCCU	
1735	GCACUGC U GGGGUCC		GGACCCC CUGAUGAG X CGAA ICAGUGC	
1742	UGGGGUC C AGGGCCC		GGGCCC CUGAUGAG X CGAA IACCCCA	
1743	GGGGUCC A GGGCCCA		UGGGCCC CUGAUGAG X CGAA IGACCCC	
1748	CCAGGGC C CACCCAG		CUGGGUG CUGAUGAG X CGAA ICCCUUG	
1749	CAGGGCC C ACCCAGU		ACUGGGU CUGAUGAG X CGAA IGGCCUG	
1750	AGGGCCC A CCCAGUG		CACUGGG CUGAUGAG X CGAA IGGCCCU	
1752	GGCCAC C CAGUGUG		CACACUG CUGAUGAG X CGAA IUGGGCC	
1753	GCCACC C AGUGUGU		ACACACU CUGAUGAG X CGAA IGUGGGC	
1754	CCCACCC A GUGUGUC		GACACAC CUGAUGAG X CGAA IGGUGGG	
1762	GUGUGUC A ACUGCAG		CUGCAGU CUGAUGAG X CGAA IACACAC	
1765	UGUCAAC U GCAGCCA		UGGCUGC CUGAUGAG X CGAA IUUGACA	
1768	CAACUGC A GCCAGUU		AACUGGC CUGAUGAG X CGAA ICAGUUG	
1771	CUGCAGC C AGUCCU		AGGAACU CUGAUGAG X CGAA ICUGCAG	
1772	UGCAGCC A GUUCUU		AAGGAAC CUGAUGAG X CGAA IGCUGCA	
1777	CCAGUUC C UUCGGGG		CCCCGAA CUGAUGAG X CGAA IAAUCUG	
1778	CAGUUC U UCGGGGC		GCCCCGA CUGAUGAG X CGAA IGAACUG	
1786	UCGGGGC C AGGAGUG		CACUCCU CUGAUGAG X CGAA ICCCCGA	
1787	CGGGGCC A GGAGUGC		GCACUCC CUGAUGAG X CGAA IGGCCCG	
1807	GGAAUGC C GAGUACU		AGUACUC CUGAUGAG X CGAA ICAUUC	
1814	CGAGUAC U GCAGGGG		CCCCUGC CUGAUGAG X CGAA IUACUCG	
1817	GUACUGC A GGGGCUC		GAGCCCC CUGAUGAG X CGAA ICAGUAC	
1823	CAGGGGC U CCCAGG		CCUGGGG CUGAUGAG X CGAA ICCCCUG	
1825	GGGGCUC C CCAGGGA		UCCUGG CUGAUGAG X CGAA IAGCCCC	
1826	GGGCUCC C CAGGGAG		CUCCUG CUGAUGAG X CGAA IGAGCCC	
1827	GGCUCCC C AGGGAGU		ACUCCU CUGAUGAG X CGAA IGGAGCC	
1828	GCUCCCC A GGGAGUA		UACUCCC CUGAUGAG X CGAA IGGGAGC	
1845	UGAAUGC C AGGCACU		AGUGCCU CUGAUGAG X CGAA ICAUUCA	
1846	GAAUGCC A GGCACUG		CAGUGCC CUGAUGAG X CGAA IGCAUUC	
1850	GCCAGGC A CUGUUUG		CAAACAG CUGAUGAG X CGAA ICCUGGC	
1852	CAGGCAC U GUUUGCC		GGCAAAC CUGAUGAG X CGAA IUGCCUG	
1859	UGUUUGC C GUGCCAC		GUGGCAC CUGAUGAG X CGAA ICAAACA	
1864	GCCGUGC C ACCCUGA		UCAGGGU CUGAUGAG X CGAA ICACGGC	
1865	CCGUGCC A CCCUGAG		CUCAGGG CUGAUGAG X CGAA IGCACGG	
1867	GUGCCAC C CUGAGUG		CACUCAG CUGAUGAG X CGAA IUGGCAC	
1868	UGCCACC C UGAGUGU		ACACUCA CUGAUGAG X CGAA IGUGGCA	
1869	GCCACCC U GAGUGUC		GACACUC CUGAUGAG X CGAA IGGUGGC	
1877	GAGUGUC A GCCCCAG		CUGGGGC CUGAUGAG X CGAA IACACUC	
1880	UGUCAGC C CCAGAAU		AUUCUGG CUGAUGAG X CGAA ICUGACA	
1881	GUCAGCC C CAGAAUG		CAUUCUG CUGAUGAG X CGAA IGCUGAC	
1882	UCAGCCC C AGAAUGG		CCAUUCU CUGAUGAG X CGAA IGGCUGA	
1883	CAGCCCC A GAAUGGC		GCCAUUC CUGAUGAG X CGAA IGGGCUG	
1891	GAAUGGC U CAGUGAC		GUCACUG CUGAUGAG X CGAA ICCAUUC	
1893	AUGGCUC A GUGACCU		AGGUCAC CUGAUGAG X CGAA IAGCCAU	
1899	CAGUGAC C UGUUUUG		CAAAACA CUGAUGAG X CGAA IUACUCG	
1900	AGUGACC U GUUUUGG		CCAAAAC CUGAUGAG X CGAA IGUCACU	

Table 34

1910	UUUGGAC C GGAGGCU		AGCCUCC CUGAUGAG X CGAA IUCCAAA	
1917	CGGAGGC U GACCAGU		ACUGGUC CUGAUGAG X CGAA ICCUCCG	
1921	GGCUGAC C AGUGUGU		ACACACU CUGAUGAG X CGAA IUCAGCC	
1922	GCUGACC A GUGUGUG		CACACAC CUGAUGAG X CGAA IGUCAGC	
1932	GUGUGGC C UGUGCCC		GGGCACA CUGAUGAG X CGAA ICCACAC	
1933	UGUGGCC U GUGCCCA		UGGGCAC CUGAUGAG X CGAA IGCCACA	
1938	CCUGUGC C CACUAUA		UAUAGUG CUGAUGAG X CGAA ICACAGG	
1939	CUGUGCC C ACUAUAA		UUAUAGU CUGAUGAG X CGAA IGCACAG	
1940	UGUGCCC A CUAUAAG		CUUAUAG CUGAUGAG X CGAA IGGCACA	
1942	UGCCCAC U AUAAGGA		UCCUUAU CUGAUGAG X CGAA IUGGGCA	
1951	UAAGGAC C CUCCCUU		AAGGGAG CUGAUGAG X CGAA IUCCUUA	
1952	AAGGACC C UCCCUUC		GAAGGGA CUGAUGAG X CGAA IGUCCUU	
1953	AGGACCC U CCCUUCU		AGAAGGG CUGAUGAG X CGAA IGGUCCU	
1955	GACCCUC C CUUCUGC		GCAGAAG CUGAUGAG X CGAA IAGGGUC	
1956	ACCCUCC C UUCUGCG		CGCAGAA CUGAUGAG X CGAA IGAGGGU	
1957	CCCUCCT U UCUGCGU		ACGCAGA CUGAUGAG X CGAA IGGAGGG	
1960	UCCCUUC U GCGUGGC		GCCACGC CUGAUGAG X CGAA IAAGGGA	
1968	GCGUGGC C CGCUGCC		GGCAGCG CUGAUGAG X CGAA ICCACGC	
1969	CGUGGCC C GCUGCCC		GGGCAGC CUGAUGAG X CGAA IGCCACG	
1972	GGCCCGC U GCCCCAG		CUGGGGC CUGAUGAG X CGAA ICGGGCC	
1975	CCGCUGC C CCAGCGG		CCGCUGG CUGAUGAG X CGAA ICAGCGG	
1976	CGCUGCC C CAGCGGU		ACCGCUG CUGAUGAG X CGAA IGCGCGG	
1977	GCUGCCC C AGCGGUG		CACCGCU CUGAUGAG X CGAA IGGCAGC	
1978	CUGCCCC A GCGGUGU		ACACCGC CUGAUGAG X CGAA IGGGCAG	
1991	GUGAAAC C UGACCUC		GAGGUCA CUGAUGAG X CGAA IUUUCAC	
1992	UGAAACC U GACCUCU		AGAGGUC CUGAUGAG X CGAA IGUUUCA	
1996	ACCUGAC C UCUCCUA		UAGGAGA CUGAUGAG X CGAA IUCAGGU	
1997	CCUGACC U CUCCUAC		GUAGGAG CUGAUGAG X CGAA IGUCAGG	
1999	UGACCUC U CCUACAU		AUGUAGG CUGAUGAG X CGAA IAGGUCA	
2001	ACCUCUC C UACAUGC		GCAUGUA CUGAUGAG X CGAA IAGAGGU	
2002	CCUCUCC U ACAUGCC		GGCAUGU CUGAUGAG X CGAA IGAGAGG	
2005	CUCCUAC A UGCCCAU		AUGGGCA CUGAUGAG X CGAA IUAGGAG	
2009	UACAUGC C CAUCUGG		CCAGAUG CUGAUGAG X CGAA ICAUGUA	
2010	ACAUGCC C AUCUGGA		UCCAGAU CUGAUGAG X CGAA IGCAUGU	
2011	CAUGCCC A UCUGGAA		UUCAGA CUGAUGAG X CGAA IGGCAUG	
2014	GCCCAUC U GGAAGUU		AACUUC CUGAUGAG X CGAA IAUGGGC	
2024	AAGUUUC C AGAUGAG		CUCAUCU CUGAUGAG X CGAA IAAACUU	
2025	AGUUUCC A GAUGAGG		CCUCAUC CUGAUGAG X CGAA IGAAACU	
2040	AGGGCGC A UGCCAGC		GCUGGCA CUGAUGAG X CGAA ICGCCCU	
2044	CGCAUGC C AGCCUUG		CAAGGCU CUGAUGAG X CGAA ICAUGCG	
2045	GCAUGCC A GCCUUGC		GCAAGGC CUGAUGAG X CGAA IGCAUGC	
2048	UGCCAGC C UUGCCCC		GGGGCAA CUGAUGAG X CGAA ICUGGCA	
2049	GCCAGCC U UGCCCCA		UGGGGCA CUGAUGAG X CGAA IGCUGGC	
2053	GCCUUGC C CCAUCAA		UUGAUGG CUGAUGAG X CGAA ICAAGGC	
2054	CCUUGCC C CAUCAAC		GUUGAUG CUGAUGAG X CGAA IGCAAGG	
2055	CUUGCCC C AUCAACU		AGUUGAU CUGAUGAG X CGAA IGGCAAG	
2056	UUGCCCC A UCAACUG		CAGUUGA CUGAUGAG X CGAA IGGGCAA	

Table 34

2059	CCCCAUC A ACUGCAC		GUGCAGU CUGAUGAG X CGAA IAUGGGG	
2062	CAUCAAC U GCACCCA		UGGGUGC CUGAUGAG X CGAA IUUGAUG	
2065	CAACUGC A CCCACUC		GAGUGGG CUGAUGAG X CGAA ICAGUUG	
2067	ACUGCAC C CACUCCU		AGGAGUG CUGAUGAG X CGAA IUGCAGU	
2068	CUGCACC C ACUCCUG		CAGGAGU CUGAUGAG X CGAA IGUGCAG	
2069	UGCACCC A CUCCUGU		ACAGGAG CUGAUGAG X CGAA IGGUGCA	
2071	CACCCAC U CCUGUGU		ACACAGG CUGAUGAG X CGAA IUGGGUG	
2073	CCCACUC C UGUGUGG		CCACACA CUGAUGAG X CGAA IAGUGGG	
2074	CCACUCC U GUGUGGA		UCCACAC CUGAUGAG X CGAA IGAGUGG	
2083	UGUGGAC C UGGAUGA		UCAUCCA CUGAUGAG X CGAA IUCCACA	
2084	GUGGACC U GGAUGAC		GUCAUCC CUGAUGAG X CGAA IUCCAC	
2092	GGAUGAC A AGGGCUG		CAGCCCU CUGAUGAG X CGAA IUCAUCC	
2098	CAAGGGC U GCCCCGC		GCGGGGC CUGAUGAG X CGAA ICCCUUG	
2101	GGGCGUC C CGCCGA		UCGGCGG CUGAUGAG X CGAA ICAGCCC	
2102	GGCUGCC C CGCCGAG		CUCGGCG CUGAUGAG X CGAA IGCAGCC	
2103	GCUGCCC C GCCGAGC		GCUCGGC CUGAUGAG X CGAA IGGCAGC	
2106	GCCCCGC C GAGCAGA		UCUGCUC CUGAUGAG X CGAA ICGGGGC	
2111	GCCGAGC A GAGAGCC		GGCUCUC CUGAUGAG X CGAA ICUCGGC	
2118	AGAGAGC C AGCCUC		GAGGGCU CUGAUGAG X CGAA ICUCUCU	
2119	GAGAGCC A GCCUCU		AGAGGGC CUGAUGAG X CGAA IGCUCUC	
2122	AGCCAGC C CUCUGAC		GUCAGAG CUGAUGAG X CGAA ICUGGCU	
2123	GCCAGCC C UCUGACG		CGUCAGA CUGAUGAG X CGAA IGCUGGC	
2124	CCAGCCC U CUGACGU		ACGUCAG CUGAUGAG X CGAA IGGCUGG	
2126	AGCCUC U GACGUCC		GGACGUC CUGAUGAG X CGAA IAGGGCU	
2133	UGACGUC C AUCAUCU		AGAUGAU CUGAUGAG X CGAA IACGUCA	
2134	GACGUCC A UCAUCUC		GAGAUGA CUGAUGAG X CGAA IGACGUC	
2137	GUCCAUC A UCUCUGC		GCAGAGA CUGAUGAG X CGAA IAUGGAC	
2140	CAUCAUC U CUGCGGU		ACCGCAG CUGAUGAG X CGAA IAUGAUG	
2142	UCAUCUC U GCGGUGG		CCACCGC CUGAUGAG X CGAA IAGAUGA	
2155	GGUUGGC A UUCUGCU		AGCAGAA CUGAUGAG X CGAA ICCAACC	
2159	GGCAUUC U GCUGGUC		GACCAGC CUGAUGAG X CGAA IAAUGCC	
2162	AUUCUGC U GGUCGUG		CACGACC CUGAUGAG X CGAA ICAGAAU	
2173	CGUGGUC U UGGGGGU		ACCCCCA CUGAUGAG X CGAA IACCACG	
2185	GGUGGUC U UUGGGAU		AUCCCAA CUGAUGAG X CGAA IACCACC	
2194	UGGGAUC C UCAUCAA		UUGAUGA CUGAUGAG X CGAA IAUCCCA	
2195	GGGAUCC U CAUCAAG		CUUGAUG CUGAUGAG X CGAA IGAUCCC	
2197	GAUCCUC A UCAAGCG		CGCUUGA CUGAUGAG X CGAA IAGGAUC	
2200	CCUCAUC A AGCGACG		CGUCGCU CUGAUGAG X CGAA IAUGAGG	
2210	CGACGGC A GCAGAAG		CUUCUGC CUGAUGAG X CGAA ICCGUCG	
2213	CGGCAGC A GAAGAUC		GAUCUUC CUGAUGAG X CGAA ICUGCCG	
2221	GAAGAUC C GGAAGUA		UACUUC CUGAUGAG X CGAA IAUCUUC	
2230	GAAGUAC A CGAUGCG		CQCAUCG CUGAUGAG X CGAA IUACUUC	
2243	CGGAGAC U GCUGCAG		CUGCAGC CUGAUGAG X CGAA IUUCUCCG	
2246	AGACUGC U GCAGGAA		UUCUUC CUGAUGAG X CGAA ICAGUCU	
2249	CUGCUGC A GGAAACG		CGUUUC CUGAUGAG X CGAA ICAGCAG	
2261	ACGGAGC U GGUGGAG		CUCCACC CUGAUGAG X CGAA ICUCCGU	
2270	GUGGAGC C GCUGACA		UGUCAGC CUGAUGAG X CGAA ICUCCAC	

Table 34

2273	GAGCCGC U GACACCU		AGGUGUC CUGAUGAG X CGAA ICGGCUC	
2277	CGCUGAC A CCUAGCG		CGCUAGG CUGAUGAG X CGAA IUCAGCG	
2279	CUGACAC C UAGCGGA		UCCGCUA CUGAUGAG X CGAA IUGUCAG	
2280	UGACACC U AGCGGAG		CUCCGCU CUGAUGAG X CGAA IGUGUCA	
2294	GCGAUGC C CAACCAG		CUGGUUG CUGAUGAG X CGAA ICAUCGC	
2295	CGAUGCC C AACCAGG		CCUGGUU CUGAUGAG X CGAA IGCAUCG	
2296	GAUGCCC A ACCAGGC		GCCUGGU CUGAUGAG X CGAA IGGCAUC	
2299	GCCCAAC C AGGCGCA		UGCGCCU CUGAUGAG X CGAA IUUGGGC	
2300	CCCAACC A GGCGCAG		CUGCGCC CUGAUGAG X CGAA IGUUGGG	
2306	CAGGCGC A GAUGCGG		CCGCAUC CUGAUGAG X CGAA ICGCCUG	
2317	GCGGAUC C UGAAAGA		UCUUUCA CUGAUGAG X CGAA IAUCCGC	
2318	CGGAUCC U GAAAGAG		CUCUUUC CUGAUGAG X CGAA IGAUCCG	
2333	ACGGAGC U GAGGAAG		CUUCCUC CUGAUGAG X CGAA ICUCCGU	
2351	AAGGUGC U UGGAUCU		AGAUGCA CUGAUGAG X CGAA ICACCUU	
2358	UUGGAUC U GGCGCUU		AAGCGCC CUGAUGAG X CGAA IAUCCAA	
2364	CUGGCGC U UUUGGCA		UGCCAAA CUGAUGAG X CGAA ICGCCAG	
2371	UUUUGGC A CAGUCUA		UAGACUG CUGAUGAG X CGAA ICCAAAA	
2373	UUGGCAC A GUCUACA		UGUAGAC CUGAUGAG X CGAA IUGCCAA	
2377	CACAGUC U ACAAGGG		CCCUUGU CUGAUGAG X CGAA IACUGUG	
2380	AGUCUAC A AGGGCAU		AUGCCCU CUGAUGAG X CGAA IUAGACU	
2386	CAAGGGC A UCUGGAU		AUCCAGA CUGAUGAG X CGAA ICCCUUG	
2389	GGGAUC U GGAUCCC		GGGAUCC CUGAUGAG X CGAA IAUGCCC	
2395	CUGGAUC C CUGAUGG		CCAUCAG CUGAUGAG X CGAA IAUCCAG	
2396	UGGAUCC C UGAUGGG		CCCAUCA CUGAUGAG X CGAA IGAUCCA	
2397	GGAUCCC U GAUGGGG		CCCCAUC CUGAUGAG X CGAA IGGAUCC	
2420	AAAAUUC C AGUGGCC		GGCCACU CUGAUGAG X CGAA IAAUUUU	
2421	AAAUUCC A GUGGCCA		UGGCCAC CUGAUGAG X CGAA IGAAUUU	
2427	CAGUGGC C AUCAAAG		CUUUGAU CUGAUGAG X CGAA ICCACUG	
2428	AGUGGCC A UCAAAGU		ACUUUGA CUGAUGAG X CGAA IGCCACU	
2431	GGCAUC A AAGUGUU		AACACUU CUGAUGAG X CGAA IAUGGCC	
2449	GGAAAAC A CAUCCCC		GGGGAUG CUGAUGAG X CGAA IUUUUCC	
2451	AAAACAC A UCCCCCA		UGGGGGA CUGAUGAG X CGAA IUGUUUU	
2454	ACACAUC C CCCAAAG		CUUUGGG CUGAUGAG X CGAA IAUGUGU	
2455	CACAUCC C CCAAAGC		GCUUUGG CUGAUGAG X CGAA IGAUGUG	
2456	ACAUCCC C CAAAGCC		GGCUUUG CUGAUGAG X CGAA IGGAUGU	
2457	CAUCCCC C AAAGCCA		UGGCUUU CUGAUGAG X CGAA IGGGAUG	
2458	AUCCCCC A AAGCCAA		UUGGCUU CUGAUGAG X CGAA IGGGGAU	
2463	CCAAAGC C AACAAAG		CUUUGUU CUGAUGAG X CGAA ICUUGGG	
2464	CAAAGCC A ACAAGA		UCUUUGU CUGAUGAG X CGAA IGCUUUG	
2467	AGCCAAC A AAGAAAU		AUUUCUU CUGAUGAG X CGAA IUUGGCU	
2476	AGAAAUC U UAGACGA		UCGUCUA CUGAUGAG X CGAA IAUUUCU	
2487	ACGAAGC A UACGUGA		UCACGUA CUGAUGAG X CGAA ICUUCGU	
2499	UGAUGGC U GGUGUGG		CCACACC CUGAUGAG X CGAA ICCAUCU	
2509	UGUGGGC U CCCCAUA		UAUGGGG CUGAUGAG X CGAA ICCACAU	
2511	UGGGCUC C CCAUAUG		CAUAUGG CUGAUGAG X CGAA IAGCCCA	
2512	GGGCUCC C CAUAUGU		ACAU AUG CUGAUGAG X CGAA IGAGCCC	
2513	GGCUCCC C AUAUGUC		GACAU AU CUGAUGAG X CGAA IGGAGCC	

Table 34

2514	GCUCCCC A UAUGUCU		AGACAU A CUGAUGAG X CGAA IGGGAGC	
2521	AUAUGUC U CCCGCCU		AGGCGGG CUGAUGAG X CGAA IACAUAU	
2523	AUGUCUC C CGCCUUC		GAAGGCG CUGAUGAG X CGAA IAGACAU	
2524	UGUCUCC C GCCUUCU		AGAAGGC CUGAUGAG X CGAA IGAGACA	
2527	CUCCCGC C UUCUGGG		CCCAGAA CUGAUGAG X CGAA ICGGGAG	
2528	UCCCGCC U UCUGGGC		GCCCAGA CUGAUGAG X CGAA ICGGGGA	
2531	CGCCUUC U GGGCAUC		GAUGCCC CUGAUGAG X CGAA IAAGGCG	
2536	UCUGGGC A UCUGCCU		AGGCAGA CUGAUGAG X CGAA ICCGAGA	
2539	GGGCAUC U GCCUGAC		GUCAGGC CUGAUGAG X CGAA IAUGCCC	
2542	CAUCUGC C UGACAUC		GAUGUCA CUGAUGAG X CGAA ICAGAUG	
2543	AUCUGCC U GACAUC		GGAUGUC CUGAUGAG X CGAA IGCAGAU	
2547	GCCUGAC A UCCACGG		CCGUGGA CUGAUGAG X CGAA IUCAGGC	
2550	UGACAUC C ACGUGGC		GCACCGU CUGAUGAG X CGAA IAUGUCA	
2551	GACAUC A CGGUGCA		UGCACCG CUGAUGAG X CGAA IGAUGUC	
2558	ACGGUGC A GCUGGUG		CACCAGC CUGAUGAG X CGAA ICACCGU	
2561	GUGCAGC U GGUGACA		UGUCACC CUGAUGAG X CGAA ICUGCAC	
2568	UGGUGAC A CAGCUUA		UAAGCUG CUGAUGAG X CGAA IUCACCA	
2570	GUGACAC A GCUUAUG		CAUAAGC CUGAUGAG X CGAA IUGUCAC	
2573	ACACAGC U UAUGCCC		GGGCAUA CUGAUGAG X CGAA ICUGUGU	
2579	CUUAUGC C CUAUGGC		GCCAUAG CUGAUGAG X CGAA ICAUAAG	
2580	UUAUGCC C UAUGGCU		AGCCAUA CUGAUGAG X CGAA IGCAUAA	
2581	UAUGCCC U AUGGCUG		CAGCCAU CUGAUGAG X CGAA IGGCAUA	
2587	CUAUGGC U GCCUCUU		AAGAGGC CUGAUGAG X CGAA ICCAUAG	
2590	UGGCUGC C UCUAAGA		UCUAAGA CUGAUGAG X CGAA ICAGCCA	
2591	GGCUGCC U CUUAGAC		GUCUAAG CUGAUGAG X CGAA IGCAGCC	
2593	CUGCCUC U UAGACCA		UGGUCUA CUGAUGAG X CGAA IAGGCAG	
2599	CUUAGAC C AUGUCCG		CGGACAU CUGAUGAG X CGAA IUCUAAG	
2600	UUAGACC A UGUCCGG		CCGGACA CUGAUGAG X CGAA IGUCUAA	
2605	CCAUGUC C GGGAAAA		UUUUCCC CUGAUGAG X CGAA ICAUUGG	
2614	GGAAAAC C GCGGACG		CGUCCGC CUGAUGAG X CGAA IUUUUCC	
2623	CGGACGC C UGGGCUC		GAGCCCA CUGAUGAG X CGAA ICGUCCG	
2624	GGACGCC U GGGCUCC		GGAGCCC CUGAUGAG X CGAA ICGGUCC	
2629	CCUGGGC U CCCAGGA		UCCUGGG CUGAUGAG X CGAA ICCCAGG	
2631	UGGGCUC C CAGGACC		GGUCCUG CUGAUGAG X CGAA IAGCCCA	
2632	GGGCUC C AGGACCU		AGGUCCU CUGAUGAG X CGAA IGAGCCC	
2633	GGCUC C A GGACCUG		CAGGUCC CUGAUGAG X CGAA IGGAGCC	
2638	CCAGGAC C UGUGAA		UUCAGCA CUGAUGAG X CGAA IUCCUGG	
2639	CAGGACC U GCUGAAC		GUUCAGC CUGAUGAG X CGAA IGUCCUG	
2642	GACCUGC U GAACTUGG		CCAGUUC CUGAUGAG X CGAA ICAGGUC	
2647	GCUGAAC U GGUGUAU		AUACACC CUGAUGAG X CGAA IUUCAGC	
2657	UGUAUGC A GAUUGCC		GGCAAUC CUGAUGAG X CGAA ICAUACA	
2664	AGAUUGC C AAGGGGA		UCCCCUU CUGAUGAG X CGAA ICAUUCU	
2665	GAUUGCC A AGGGGAU		AUCCCCU CUGAUGAG X CGAA IGCAAUC	
2677	GAUGAGC U ACCUGGA		UCCAGGU CUGAUGAG X CGAA ICUCAUC	
2680	GAGCUAC C UGGAGGA		UCCUCCA CUGAUGAG X CGAA IUAGCUC	
2681	AGCUACC U GGAGGAU		AUCCUCC CUGAUGAG X CGAA IGUAGCU	
2696	GUGCGGC U CGUACAC		GUGUACG CUGAUGAG X CGAA ICCGCAC	

Table 34

2702	CUCGUAC A CAGGGAC		GUCCUG CUGAUGAG X CGAA IUACGAG	
2704	CGUACAC A GGGACUU		AAGUCCC CUGAUGAG X CGAA IUGUACG	
2710	CAGGGAC U UGGCCGC		GCGGCCA CUGAUGAG X CGAA IUCCUG	
2715	ACUUGGC C GCUCGGA		UCCGAGC CUGAUGAG X CGAA ICCAAGU	
2718	UGGCCGC U CGGAACG		CGUCCG CUGAUGAG X CGAA ICGGCCA	
2729	AACGUGC U GGUCAAG		CUUGACC CUGAUGAG X CGAA ICACGUU	
2734	GCUGGUC A AGAGUCC		GGACUCU CUGAUGAG X CGAA IACCAGC	
2741	AAGAGUC C CAACCAU		AUGGUUG CUGAUGAG X CGAA IACUCUU	
2742	AGAGUCC C AACCAUG		CAUGGUU CUGAUGAG X CGAA IGACUCU	
2743	GAGUCCC A ACCAUGU		ACAUGGU CUGAUGAG X CGAA IGGACUC	
2746	UCCCAAC C AUGUCAA		UUGACAU CUGAUGAG X CGAA IUUGGGA	
2747	CCCAACC A UGUCAAA		UUUGACA CUGAUGAG X CGAA IGUUGGG	
2752	CCAUGUC A AAAUUAC		GUAUUU CUGAUGAG X CGAA IACAUGG	
2760	AAAUUAC A GACUUCG		CGAAGUC CUGAUGAG X CGAA IUAAUUU	
2764	UACAGAC U UCGGGCU		AGCCCGA CUGAUGAG X CGAA IUCUGUA	
2771	UUCGGGC U GGCUCGG		CCGAGCC CUGAUGAG X CGAA ICCCGAA	
2775	GGCUGGC U CGGCGC		GCAGCCG CUGAUGAG X CGAA ICCAGCC	
2780	GCUCGGC U GCUGGAC		GUCCAGC CUGAUGAG X CGAA ICCGAGC	
2783	CGGCGC U GGACAUU		AAUGUCC CUGAUGAG X CGAA ICAGCCG	
2788	GCUGGAC A UUGACGA		UCGUCAA CUGAUGAG X CGAA IUCCAGC	
2799	ACGAGAC A GAGUACC		GGUACUC CUGAUGAG X CGAA IUCUCGU	
2806	AGAGUAC C AUGCAGA		UCUGCAU CUGAUGAG X CGAA IUACUCU	
2807	GAGUACC A UGCAGAU		AUCUGCA CUGAUGAG X CGAA IGUACUC	
2811	ACCAUGC A GAUGGGG		CCCAUC CUGAUGAG X CGAA ICAUGGU	
2821	UGGGGGC A AGGUGCC		GGCACCU CUGAUGAG X CGAA ICCCCCA	
2828	AAGGUGC C CAUCAAG		CUUGAUG CUGAUGAG X CGAA ICACCUU	
2829	AGGUGCC C AUCAAGU		ACUUGAU CUGAUGAG X CGAA IGCACCU	
2830	GGUGCCC A UCAAGUG		CACUUGA CUGAUGAG X CGAA IGGACC	
2833	GCCCAUC A AGUGGAU		AUCCACU CUGAUGAG X CGAA IAUGGGC	
2846	AUGGCGC U GGAGUCC		GGACUCC CUGAUGAG X CGAA ICGCCAU	
2853	UGGAGUC C AUUCUCC		GGAGAAU CUGAUGAG X CGAA IACUCCA	
2854	GGAGUCC A UUCUCCG		CGGAGAA CUGAUGAG X CGAA IGACUCC	
2858	UCCAUC U CCGCCG		CCGGCGG CUGAUGAG X CGAA IAAUGGA	
2860	CAUUCUC C GCCGGCG		CGCCGGC CUGAUGAG X CGAA IAGAAUG	
2863	UCUCCGC C GCGGUU		AACCGCC CUGAUGAG X CGAA ICGGAGA	
2872	GCGGUUC A CCCACCA		UGGUGGG CUGAUGAG X CGAA IAACCGC	
2874	GGUUCAC C CACCAGA		UCUGGUG CUGAUGAG X CGAA IUGAACC	
2875	GUUCACC C ACCAGAG		CUCUGGU CUGAUGAG X CGAA IGUGAAC	
2876	IUCACCC A CCAGAGU		ACUCUGG CUGAUGAG X CGAA IGGUGAA	
2878	CACCCAC C AGAGUGA		UCACUCU CUGAUGAG X CGAA IUUGGUG	
2879	ACCCACC A GAGUGAU		AUCACUC CUGAUGAG X CGAA IGUGGGU	
2907	GUGUGAC U GUGUGGG		CCCACAC CUGAUGAG X CGAA IUCACAC	
2918	UGGGAGC U GAUGACU		AGUCAUC CUGAUGAG X CGAA ICUCCCA	
2925	UGAUGAC U UUUGGGG		CCCCAAA CUGAUGAG X CGAA IUCAUCA	
2934	UUUGGGC C AAACCUU		AAGGUUU CUGAUGAG X CGAA ICCCCAA	
2935	UGGGGCC A AACCUUA		UAAGGUU CUGAUGAG X CGAA IGGCCCA	
2939	GCCAAAC C UUACGAU		AUCGUAA CUGAUGAG X CGAA IUUGGC	

Table 34

2940	CCAAACC U UACGAUG		CAUCGUA CUGAUGAG X CGAA IGUUUGG	
2953	UGGGAUC C CAGCCCCG		CGGGCUG CUGAUGAG X CGAA IAUCCCA	
2954	GGGAUCC C AGCCCCG		CCGGGCU CUGAUGAG X CGAA IGAUCC	
2955	GGAUCCC A GCCCCGG		CCCGGGC CUGAUGAG X CGAA IGGAUCC	
2958	UCCCAGC C CGGGAGA		UCUCCCG CUGAUGAG X CGAA ICUGGGA	
2959	CCCAGCC C GGGAGAU		AUCUCCC CUGAUGAG X CGAA IGCUGGG	
2968	GGAGAUC C CUGACCU		AGGUCAG CUGAUGAG X CGAA IAUUCC	
2969	GAGAUCC C UGACCUG		CAGGUCA CUGAUGAG X CGAA IGAUCUC	
2970	AGAUCCC U GACCUGC		GCAGGUC CUGAUGAG X CGAA IGGAUCU	
2974	CCCUGAC C UGCUGGA		UCCAGCA CUGAUGAG X CGAA IUCAGGG	
2975	CCUGACC U GCUGGAA		UUCCAGC CUGAUGAG X CGAA IGUCAGG	
2978	GACCUGC U GGAAAAG		CUUUUCC CUGAUGAG X CGAA ICAGGUC	
2996	GAGCGGC U GCCCCAG		CUGGGGC CUGAUGAG X CGAA ICCGCUC	
2999	CGGCGUC C CCAGCCC		GGGCGUG CUGAUGAG X CGAA ICAGCCG	
3000	GGCUGCC C CAGCCCC		GGGGCUG CUGAUGAG X CGAA IGCAGCC	
3001	GCUGCCC C AGCCCCC		GGGGGCU CUGAUGAG X CGAA IGGCAGC	
3002	CUGCCCC A GCCCCC		GGGGGGC CUGAUGAG X CGAA IGGGCAG	
3005	CCCCAGC C CCCCAUC		GAUGGGG CUGAUGAG X CGAA ICUGGGG	
3006	CCCAGCC C CCCAUCU		AGAUGGG CUGAUGAG X CGAA IGCUGGG	
3007	CCAGCCC C CCAUCUG		CAGAUGG CUGAUGAG X CGAA IGGCUGG	
3008	CAGCCCC C CAUCUGC		GCAGAUG CUGAUGAG X CGAA IGGGCUG	
3009	AGCCCCC C AUCUGCA		UGCAGAU CUGAUGAG X CGAA IGGGGCU	
3010	GCCCCC A UCUGCAC		GUGCAGA CUGAUGAG X CGAA IGGGGGC	
3013	CCCAUC U GCACCAU		AUGGUGC CUGAUGAG X CGAA IAUGGGG	
3016	CAUCUGC A CCAUUGA		UCAAUGG CUGAUGAG X CGAA ICAGAUG	
3018	UCUGCAC C AUUGAUG		CAUCAAU CUGAUGAG X CGAA IUGCAGA	
3019	CUGCACC A UUGAUGU		ACAUCAA CUGAUGAG X CGAA IGUGCAG	
3028	UGAUGUC U ACAUGAU		AUCAUGU CUGAUGAG X CGAA IACAUCA	
3031	UGUCUAC A UGAUCAU		AUGAUCA CUGAUGAG X CGAA IUAGACA	
3037	CAUGAUC A UGGUCAA		UUGACCA CUGAUGAG X CGAA IAUCAUG	
3043	CAUGGUC A AAUGUUG		CAACAUC CUGAUGAG X CGAA IACCAUG	
3061	GAUUGAC U CUGAAUG		CAUUCAG CUGAUGAG X CGAA IUCAAUC	
3063	UUGACUC U GAAUGUC		GACAUUC CUGAUGAG X CGAA IAGUCAA	
3074	UGUCGGC C AAGAUUC		GAAUCUU CUGAUGAG X CGAA ICCGACA	
3075	GUCGGCC A AGAUUCC		GGAUUCU CUGAUGAG X CGAA IGCCGAC	
3082	AAGAUUC C GGGAGUU		AACUCCC CUGAUGAG X CGAA IAAUCUU	
3096	UGGUGUC U GAAUUCU		AGAAUUC CUGAUGAG X CGAA IACACCA	
3103	UGAAUUC U CCCGCAU		AUGCGGG CUGAUGAG X CGAA IAAUUCA	
3105	AAUUCUC C CGCAUGG		CCAUGCG CUGAUGAG X CGAA IAGAAUU	
3106	AUUCUCC C GCAUGGC		GCCAUGC CUGAUGAG X CGAA IGAGAAU	
3109	CUCCCGC A UGGCCAG		CUGGCCA CUGAUGAG X CGAA ICGGGAG	
3114	GCAUGGC C AGGGACC		GGUCCCU CUGAUGAG X CGAA ICCAUGC	
3115	CAUGGCC A GGGACCC		GGGUCCC CUGAUGAG X CGAA IGCCAUG	
3121	CAGGGAC C CCCAGCG		CGCUGGG CUGAUGAG X CGAA IUCCUG	
3122	AGGGACC C CCAGCGC		GCGCUGG CUGAUGAG X CGAA IGUCCCU	
3123	GGGACCC C CAGCGCU		AGCGCUG CUGAUGAG X CGAA IGGUCCC	
3124	GGACCCC C AGCGCUU		AAGCGCU CUGAUGAG X CGAA IGGGUCC	

Table 34

3125	GACCCCC A GCGCUUU		AAAGCGC CUGAUGAG X CGAA IGGGGUC	
3130	CCAGCGC U UUGUGGU		ACCACAA CUGAUGAG X CGAA ICGUGG	
3139	UGGGGUC A UCCAGAA		UUCUGGA CUGAUGAG X CGAA IACCACA	
3142	GGUCAUC C AGAAUGA		UCAUUCU CUGAUGAG X CGAA IAUGACC	
3143	GUCAUCC A GAAUGAG		CUCAUUC CUGAUGAG X CGAA IGAUGAC	
3154	UGAGGAC U UGGGCC		GGGCCCA CUGAUGAG X CGAA IUCCUCA	
3160	CUUGGGC C CAGCCAG		CUGGCUG CUGAUGAG X CGAA ICCCAAG	
3161	UUGGGCC C AGCCAGU		ACUGGCU CUGAUGAG X CGAA IGCCCAA	
3162	UGGGCCC A GCCAGUC		GACUGGC CUGAUGAG X CGAA IGGCCCA	
3165	GCCAGC C AGUCCCU		AGGGACU CUGAUGAG X CGAA ICUGGGC	
3166	CCCAGCC A GUCCCUU		AAGGGAC CUGAUGAG X CGAA IGCUGGG	
3170	GCCAGUC C CUUGGAC		GUCCAAG CUGAUGAG X CGAA IACUGGC	
3171	CCAGUCC C UUGGACA		UGUCCAA CUGAUGAG X CGAA IGACUGG	
3172	CAGUCCC U UGGACAG		CUGUCCA CUGAUGAG X CGAA IGGACUG	
3178	CUUGGAC A GCACCUU		AAGGUGC CUGAUGAG X CGAA IUCCAAG	
3181	GGACAGC A CCUUCUA		UAGAAGG CUGAUGAG X CGAA ICUGUCC	
3183	ACAGCAC C UUCUACC		GGUAGAA CUGAUGAG X CGAA IUGCUGU	
3184	CAGCACC U UCUACCG		CGGUAGA CUGAUGAG X CGAA IGUGCUG	
3187	CACCUUC U ACCGCUC		GAGCGGU CUGAUGAG X CGAA IAAGGUG	
3190	CUUCUAC C GCUCACU		AGUGAGC CUGAUGAG X CGAA IUAGAAG	
3193	CUACCGC U CACUGCU		AGCAGUG CUGAUGAG X CGAA ICGGUAG	
3195	ACCGCUC A CUGCUGG		CCAGCAG CUGAUGAG X CGAA IAGCGGU	
3197	CGCUCAC U GCUGGAG		CUCCAGC CUGAUGAG X CGAA IUGAGCG	
3200	UCACUGC U GGAGGAC		GUCCUCC CUGAUGAG X CGAA ICAGUGA	
3214	CGAUGAC A UGGGGGA		UCCCCCA CUGAUGAG X CGAA IUCAUCG	
3223	GGGGGAC C UGUGGGA		UCCACCA CUGAUGAG X CGAA IUCCCCC	
3224	GGGGACC U GGUGGAU		AUCCACC CUGAUGAG X CGAA IGUCCCC	
3234	UGGAUGC U GAGGAGU		ACUCCUC CUGAUGAG X CGAA ICAUCCA	
3245	GAGUAUC U GGUACCC		GGGUACC CUGAUGAG X CGAA IAUACUC	
3251	CUGGUAC C CCAGCAG		CUGCUGG CUGAUGAG X CGAA IUACCAG	
3252	UGGUACC C CAGCAGG		CCUGCUG CUGAUGAG X CGAA IGUACCA	
3253	GGUACCC C AGCAGGG		CCCUGCU CUGAUGAG X CGAA IGGUACC	
3254	GUACCCC A GCAGGGC		GCCCUGC CUGAUGAG X CGAA IGGGUAC	
3257	CCCCAGC A GGGCUUC		GAAGCCC CUGAUGAG X CGAA ICUGGGG	
3262	GCAGGGC U UCUCUCG		CAGAAGA CUGAUGAG X CGAA ICCUCGC	
3265	GGGCUUC U UCUGUCC		GGACAGA CUGAUGAG X CGAA IAAGCCC	
3268	CUUCUUC U GUCCAGA		UCUGGAC CUGAUGAG X CGAA IAAGAAG	
3272	UUCUGUC C AGACCCU		AGGGUCU CUGAUGAG X CGAA IACAGAA	
3273	UCUGUCC A GACCCUG		CAGGGUC CUGAUGAG X CGAA IGACAGA	
3277	UCCAGAC C CUGCCCC		GGGGCAG CUGAUGAG X CGAA IUCUGGA	
3278	CCAGACC C UGCCCCG		CGGGGCA CUGAUGAG X CGAA IGUCUGG	
3279	CAGACCC U GCCCCCG		CCGGGGC CUGAUGAG X CGAA IGGUCUG	
3282	ACCCUGC C CCGGGCG		CGCCCCG CUGAUGAG X CGAA ICAGGGU	
3283	CCCUGCC C CGGGCGC		GCGCCCC CUGAUGAG X CGAA IGCAGGG	
3284	CCUGCCC C GGGCGCU		AGCGCCC CUGAUGAG X CGAA IGGCAGG	
3291	CGGGCGC U GGGGGCA		UGCCCCC CUGAUGAG X CGAA ICGCCCG	
3298	UGGGGGC A UGUCCA		UGGACCA CUGAUGAG X CGAA ICCCCCA	

Table 34

3304	CAUGGUC C ACCACAG		CUGUGGU CUGAUGAG X CGAA IACCAUG	
3305	AUGGUCC A CCACAGG		CCUGUGG CUGAUGAG X CGAA IGACCAU	
3307	GGUCCAC C ACAGGCA		UGCCUGU CUGAUGAG X CGAA IUGGACC	
3308	GUCCACC A CAGGCAC		GUGCCUG CUGAUGAG X CGAA IGUGGAC	
3310	CCACCAC A GGCACCG		CGGUGCC CUGAUGAG X CGAA IUGGUGG	
3314	CACAGGC A CCGCAGC		GCUGCGG CUGAUGAG X CGAA ICCUGUG	
3316	CAGGCAC C GCAGCUC		GAGCUGC CUGAUGAG X CGAA IUGCCUG	
3319	GCACGC A GCUCAUC		GAUGAGC CUGAUGAG X CGAA ICGGUGC	
3322	CCGCAGC U CAUCUAC		GUAGAUG CUGAUGAG X CGAA ICUGCGG	
3324	GCAGCUC A UCUACCA		UGGUAGA CUGAUGAG X CGAA IAGCUGC	
3327	GCUCAUC U ACCAGGA		UCCUGGU CUGAUGAG X CGAA IAUGAGC	
3330	CAUCUAC C AGGAGUG		CACUCCU CUGAUGAG X CGAA IUAGAUG	
3331	AUCUACC A GGAGUGG		CCACUCC CUGAUGAG X CGAA IGUAGAU	
3349	UGGGGAC C UGACACU		AGUGUCA CUGAUGAG X CGAA IUCCCCA	
3350	GGGGACC U GACACUA		UAGUGUC CUGAUGAG X CGAA IGUCCCC	
3354	ACCUGAC A CUAGGGC		GCCCUAG CUGAUGAG X CGAA IUCAGGU	
3356	CUGACAC U AGGGCUG		CAGCCCU CUGAUGAG X CGAA IUGUCAG	
3362	CUAGGGC U GGAGCCC		GGGCUC CUGAUGAG X CGAA ICCCUAG	
3368	CUGGAGC C CUCUGAA		UUCAGAG CUGAUGAG X CGAA ICUCCAG	
3369	UGGAGCC C UCUGAAG		CUUCAGA CUGAUGAG X CGAA IGCUCCA	
3370	GGAGCCC U CUGAAGA		UCUUCAG CUGAUGAG X CGAA IGGCUCC	
3372	AGCCUC U GAAGAGG		CCUCUUC CUGAUGAG X CGAA IAGGGCU	
3384	AGGAGGC C CCCAGGU		ACCUGGG CUGAUGAG X CGAA ICCUCCU	
3385	GGAGGCC C CCAGGUC		GACCUGG CUGAUGAG X CGAA IGGCUCC	
3386	GAGGCC C CAGGUCU		AGACCUG CUGAUGAG X CGAA IGGCCUC	
3387	AGGCCCC C AGGUCUC		GAGACCU CUGAUGAG X CGAA IGGGCCU	
3388	GGCCCC A GGUCUCC		GGAGACC CUGAUGAG X CGAA IGGGGCC	
3393	CCAGGUC U CCACUGG		CCAGUGG CUGAUGAG X CGAA IACCUGG	
3395	AGGUCUC C ACUGGCA		UGCCAGU CUGAUGAG X CGAA IAGACCU	
3396	GGUCUCC A CUGGCAC		GUGCCAG CUGAUGAG X CGAA IGAGACC	
3398	UCUCCAC U GGCACCC		GGGUGCC CUGAUGAG X CGAA IUGGAGA	
3402	CACUGGC A CCCUCCG		CGGAGGG CUGAUGAG X CGAA ICCAGUG	
3404	CUGGCAC C CUCCGAA		UUCGGAG CUGAUGAG X CGAA IUGCCAG	
3405	UGGCACC C UCCGAAG		CUUCGGA CUGAUGAG X CGAA IGUGCCA	
3406	GGCACCC U CCGAAGG		CCUUCGG CUGAUGAG X CGAA IGGUGCC	
3408	CACCCUC C GAAGGGG		CCCUUUC CUGAUGAG X CGAA IAGGGUG	
3417	AAGGGGC U GGCUCCG		CGGAGCC CUGAUGAG X CGAA ICCCUU	
3421	GGCUGGC U CCGAUGU		ACAUCGG CUGAUGAG X CGAA ICCAGCC	
3423	CUGGCUC C GAUGUAU		AUACAUC CUGAUGAG X CGAA IAGCCAG	
3442	UGGUGAC C UGGGAU		AUUCCA CUGAUGAG X CGAA IUCACCA	
3443	GGUGACC U GGGAAUG		CAUCCC CUGAUGAG X CGAA IGUCACC	
3456	UGGGGGC A GCCAAGG		CCUUGGC CUGAUGAG X CGAA ICCCCCA	
3459	GGGCAGC C AAGGGGC		GCCCUU CUGAUGAG X CGAA ICUGCCC	
3460	GGCAGCC A AGGGGCU		AGCCCU CUGAUGAG X CGAA IGCUGCC	
3467	AAGGGGC U GCAAAGC		GCUUUGC CUGAUGAG X CGAA ICCCUU	
3470	GGGCUGC A AAGCCUC		GAGGCUU CUGAUGAG X CGAA ICAGCCC	
3475	GCAAAGC C UCCCCAC		GUGGGGA CUGAUGAG X CGAA ICUUGC	

Table 34

3476	CAAAGCC U CCCACA		UGUGGG CUGAUGAG X CGAA IGCUUUG	
3478	AAGCCUC C CCACACA		UGUGUG CUGAUGAG X CGAA IAGGCUU	
3479	AGCCUCC C CACACAU		AUGUGUG CUGAUGAG X CGAA IGAGGCU	
3480	GCCUCCC C ACACAUG		CAUGUGU CUGAUGAG X CGAA IGGAGGC	
3481	CCUCCCC A CACAUGA		UCAUGUG CUGAUGAG X CGAA IGGGAGG	
3483	UCCCCAC A CAUGACC		GGUCAUG CUGAUGAG X CGAA IUGGGGA	
3485	CCCACAC A UGACCCC		GGGUGA CUGAUGAG X CGAA IUGUGGG	
3490	ACAUGAC C CCAGCCC		GGGUGG CUGAUGAG X CGAA IUCAUGU	
3491	CAUGACC C CAGCCCU		AGGGCUG CUGAUGAG X CGAA IGUCAUG	
3492	AUGACCC C AGCCUCU		GAGGGCU CUGAUGAG X CGAA IGGUCAU	
3493	UGACCCC A GCCUCU		AGAGGGC CUGAUGAG X CGAA IGGGUCA	
3496	CCCCAGC C CUCUACA		UGUAGAG CUGAUGAG X CGAA ICUGGGG	
3497	CCCAGCC C UCUCAG		CUGUAGA CUGAUGAG X CGAA IGCUGGG	
3498	CCAGCCC U CUACAGC		GCUGUAG CUGAUGAG X CGAA IGGCUGG	
3500	AGCCUCU U ACAGCGG		CCGCUGU CUGAUGAG X CGAA IAGGGCU	
3503	CCUCUAC A GCGGUAC		GUACCGC CUGAUGAG X CGAA IUAGAGG	
3511	GCGGUAC A GUGAGGA		UCCUCAC CUGAUGAG X CGAA IUACCGC	
3520	UGAGGAC C CCACAGU		ACUGUGG CUGAUGAG X CGAA IUCCUCA	
3521	GAGGACC C CACAGUA		UACUGUG CUGAUGAG X CGAA IGUCCUC	
3522	AGGACCC C ACAGUAC		GUACUGU CUGAUGAG X CGAA IGGUCCU	
3523	GGACCCC A CAGUACC		GGUACUG CUGAUGAG X CGAA IGGGUCC	
3525	ACCCAC A GUACCCC		GGGGUAC CUGAUGAG X CGAA IUGGGGU	
3530	ACAGUAC C CCUGCCC		GGGCAGG CUGAUGAG X CGAA IUACUGU	
3531	CAGUACC C CUGCCCU		AGGGCAG CUGAUGAG X CGAA IGUACUG	
3532	AGUACCC C UGCCCUC		GAGGGCA CUGAUGAG X CGAA IGGUACU	
3533	GUACCCC U GCCUCU		AGAGGGC CUGAUGAG X CGAA IGGGUAC	
3536	CCCUGC C CUCUGAG		CUCAGAG CUGAUGAG X CGAA ICAGGGG	
3537	CCCUGCC C UCUGAGA		UCUCAGA CUGAUGAG X CGAA IGCAGGG	
3538	CCUGCCC U CUGAGAC		GUCUCAG CUGAUGAG X CGAA IGGCAGG	
3540	UGCCCUC U GAGACUG		CAGUCUC CUGAUGAG X CGAA IAGGGCA	
3546	CUGAGAC U GAUGGCU		AGCCAUC CUGAUGAG X CGAA IUCUCAG	
3553	UGAUGGC U ACGUUGC		GCAACGU CUGAUGAG X CGAA ICCAUCA	
3561	ACGUUGC C CCCUGA		UCAGGGG CUGAUGAG X CGAA ICAACGU	
3562	CGUUGCC C CCCUGAC		GUCAGGG CUGAUGAG X CGAA IGCAACG	
3563	GUUGCCC C CCUGACC		GGUCAGG CUGAUGAG X CGAA IGGCAAC	
3564	UUGCCCC C CUGACCU		AGGUCAG CUGAUGAG X CGAA IGGGCAA	
3565	UGCCCCC C UGACCU		CAGGUCA CUGAUGAG X CGAA IGGGGCA	
3566	GCCCCC U GACCUGC		GCAGGUC CUGAUGAG X CGAA IGGGGGC	
3570	CCCUGAC C UGCAGCC		GGCUGCA CUGAUGAG X CGAA IUCAGGG	
3571	CCUGACC U GCAGCCC		GGGCUGC CUGAUGAG X CGAA IGUCAGG	
3574	GACCUGC A GCCCCA		UGGGGGC CUGAUGAG X CGAA ICAGGUC	
3577	CUGCAGC C CCCAGCC		GGCUGGG CUGAUGAG X CGAA ICUGCAG	
3578	UGCAGCC C CCAGCCU		AGGCUGG CUGAUGAG X CGAA IGCUGCA	
3579	GCAGCCC C CAGCCUG		CAGGCUG CUGAUGAG X CGAA IGGCUGC	
3580	CAGCCCC C AGCCUGA		UCAGGCU CUGAUGAG X CGAA IGGGCUG	
3581	AGCCCCC A GCCUGAA		UUCAGGC CUGAUGAG X CGAA IGGGGCU	
3584	CCCCAGC C UGAUAU		AUAUUA CUGAUGAG X CGAA ICUGGGG	

Table 34

3585	CCCAGCC U GAAUAUG		CAUAUUC CUGAUGAG X CGAA IGCUGGG	
3598	UGUGAAC C AGCCAGA		UCUGGCU CUGAUGAG X CGAA IUUCACA	
3599	GUGAACC A GCCAGAU		AUCUGGC CUGAUGAG X CGAA IGUUCAC	
3602	AACCAGC C AGAUGUU		AACAUCU CUGAUGAG X CGAA ICUGGUU	
3603	ACCAGCC A GAUGUUC		GAACAUC CUGAUGAG X CGAA IGCUGGU	
3614	GUUCGGC C CCAGCCC		GGGCUUG CUGAUGAG X CGAA ICCGAAC	
3615	UUCGGCC C CAGCCCC		GGGGCUG CUGAUGAG X CGAA IGCCGAA	
3616	UCGGCCC C AGCCCCC		GGGGGCU CUGAUGAG X CGAA IGGCCGA	
3617	CGGCCCC A GCCCCCU		AGGGGGC CUGAUGAG X CGAA IGGGCCG	
3620	CCCCAGC C CCCUUCG		CGAAGGG CUGAUGAG X CGAA ICUGGGG	
3621	CCCAGCC C CCUUCGC		GCGAAGG CUGAUGAG X CGAA IGCUGGG	
3622	CCAGCCC C CUUCGCC		GGCGAAG CUGAUGAG X CGAA IGGCUGG	
3623	CAGCCCC C UUCGCCC		GGGCGAA CUGAUGAG X CGAA IGGGCUG	
3624	AGCCCCC U UCGCCCC		GGGGCGA CUGAUGAG X CGAA IGGGGCU	
3629	CCUUCGC C CCGAGAG		CUCUCGG CUGAUGAG X CGAA ICGAAGG	
3630	CUUCGCC C CGAGAGG		CCUCUCG CUGAUGAG X CGAA ICGGAAG	
3631	UUCGCCC C GAGAGGG		CCCUCUC CUGAUGAG X CGAA IGGCGAA	
3640	AGAGGGC C CUCUGCC		GGCAGAG CUGAUGAG X CGAA ICCUCUC	
3641	GAGGGCC C UCUGCCU		AGGCAGA CUGAUGAG X CGAA IGCCUC	
3642	AGGGCCC U CUGCCUG		CAGGCAG CUGAUGAG X CGAA IGGCCCU	
3644	GGCCUCU U GCCUGCU		AGCAGGC CUGAUGAG X CGAA IAGGGCC	
3647	CCUCUGC C UGCUGCC		GGCAGCA CUGAUGAG X CGAA ICAGAGG	
3648	CUCUGCC U GCUGCCC		GGGCAGC CUGAUGAG X CGAA IGCAGAG	
3651	UGCCUGC U GCCCGAC		GUCGGGC CUGAUGAG X CGAA ICAGGCA	
3654	CUGCUGC C CGACCUG		CAGGUCG CUGAUGAG X CGAA ICAGCAG	
3655	UGCUGCC C GACCUGC		GCAGGUC CUGAUGAG X CGAA IGCAGCA	
3659	GCCCGAC C UGCUGGU		ACCAGCA CUGAUGAG X CGAA IUCGGGC	
3660	CCCGACC U GCUGGUG		CACCAGC CUGAUGAG X CGAA IGUCGGG	
3663	GACCUGC U GGUGCCA		UGGCACC CUGAUGAG X CGAA ICAGGUC	
3669	CUGGUGC C ACUCUGG		CCAGAGU CUGAUGAG X CGAA ICACCAG	
3670	UGGUGCC A CUCUGGA		UCCAGAG CUGAUGAG X CGAA IGCACCA	
3672	GUGCCAC U CUGGAAA		UUUCCAG CUGAUGAG X CGAA IUUGCAC	
3674	GCCACUC U GGAAAGG		CCUUUCC CUGAUGAG X CGAA IAGUGGC	
3683	GAAAGGC C CAAGACU		AGUCUUG CUGAUGAG X CGAA ICCUUUC	
3684	AAAGGCC C AAGACUC		GAGUCUU CUGAUGAG X CGAA IGCCUUU	
3685	AAGGCCC A AGACUCU		AGAGUCU CUGAUGAG X CGAA IGGCCUU	
3690	CCAAGAC U CUCUCCC		GGGAGAG CUGAUGAG X CGAA IUCUUGG	
3692	AAGACUC U CUCCCCA		UGGGGAG CUGAUGAG X CGAA IAGUCUU	
3694	GACUCUC U CCCCAGG		CCUGGGG CUGAUGAG X CGAA IAGAGUC	
3696	CUCUCUC C CCAGGGA		UCCCUGG CUGAUGAG X CGAA IAGAGAG	
3697	UCUCUCC C CAGGGAA		UUCCCUG CUGAUGAG X CGAA IGAGAGA	
3698	CUCUCCC C AGGGAAG		CUUCCCU CUGAUGAG X CGAA IGGAGAG	
3699	UCUCCCC A GGAAGA		UCUCCC CUGAUGAG X CGAA IGGGAGA	
3718	GGUCGUC A AAGACGU		ACGUCUU CUGAUGAG X CGAA IACGACC	
3732	UUUUUGC C UUUGGGG		CCCCAAA CUGAUGAG X CGAA ICAAAAA	
3733	UUUUGCC U UUGGGGG		CCCCCAA CUGAUGAG X CGAA IGCAAAA	
3744	GGGGUGC C GUGGAGA		UCUCCAC CUGAUGAG X CGAA ICACCCC	

Table 34

3754	GGAGAAC C CCGAGUA		UACUCGG CUGAUGAG X CGAA IUUCUCC	
3755	GAGAACC C CGAGUAC		GUACUCG CUGAUGAG X CGAA IGUUCUC	
3756	AGAACCC C GAGUACU		AGUACUC CUGAUGAG X CGAA IGGUUCU	
3763	CGAGUAC U UGACACC		GGUGUCA CUGAUGAG X CGAA IUACUCG	
3768	ACUUGAC A CCCCAGG		CCUGGGG CUGAUGAG X CGAA IUCAAGU	
3770	UUGACAC C CCAGGGA		UCCUCGG CUGAUGAG X CGAA IUGUCAA	
3771	UGACACC C CAGGGAG		CUCCUCG CUGAUGAG X CGAA IGUGUCA	
3772	GACACCC C AGGGAGG		CCUCCCU CUGAUGAG X CGAA IGGUGUC	
3773	ACACCCC A GGGAGGA		UCCUCCC CUGAUGAG X CGAA IGGGUGU	
3783	GAGGAGC U GCCCCUC		GAGGGGC CUGAUGAG X CGAA ICUCCUC	
3786	GAGCUGC C CCUCAGC		GCUGAGG CUGAUGAG X CGAA ICAGCUC	
3787	AGCUGCC C CUCAGCC		GGCUGAG CUGAUGAG X CGAA IGCAGCU	
3788	GCUGCCC C UCAGCCC		GGGCUGA CUGAUGAG X CGAA IGGCAGC	
3789	CUGCCCC U CAGCCCC		GGGGCUG CUGAUGAG X CGAA IGGGCAG	
3791	GCCCCUC A GCCCCAC		GUGGGGC CUGAUGAG X CGAA IAGGGGC	
3794	CCUCAGC C CCACCCU		AGGGUGG CUGAUGAG X CGAA ICUGAGG	
3795	CUCAGCC C CACCCUC		GAGGGUG CUGAUGAG X CGAA IGCUGAG	
3796	UCAGCCC C ACCUCC		GGAGGGU CUGAUGAG X CGAA IGGCUGA	
3797	CAGCCCC A CCUCCU		AGGAGGG CUGAUGAG X CGAA IGGGCUG	
3799	GCCCCAC C CUCCUCC		GGAGGAG CUGAUGAG X CGAA IUGGGGC	
3800	CCCCACC C UCCUCCU		AGGAGGA CUGAUGAG X CGAA IGUGGGG	
3801	CCCACCC U CCUCCUG		CAGGAGG CUGAUGAG X CGAA IGGUGGG	
3803	CACCCUC C UCCUGCC		GGCAGGA CUGAUGAG X CGAA IAGGGUG	
3804	ACCCUCC U CCUGCCU		AGGCAGG CUGAUGAG X CGAA IGAGGGU	
3806	CCUCCUC C UGCCUUC		GAAGGCA CUGAUGAG X CGAA IAGGAGG	
3807	CUCCUCC U GCCUUA		UGAAGGC CUGAUGAG X CGAA IGAGGAG	
3810	CUCCUGC C UUCAGCC		GGCUGAA CUGAUGAG X CGAA ICAGGAG	
3811	UCCUGCC U UCAGCCC		GGGCUGA CUGAUGAG X CGAA IGCAGGA	
3814	UGCCUUC A GCCCAGC		GCUGGGC CUGAUGAG X CGAA IAAGGCA	
3817	CUUCAGC C CAGCCU		AAGGCUG CUGAUGAG X CGAA ICUGAAG	
3818	UUCAGCC C AGCCUUC		GAAGGCU CUGAUGAG X CGAA IGCUGAA	
3819	UCAGCCC A GCCUUCG		CGAAGGC CUGAUGAG X CGAA IGGCUGA	
3822	GCCCAGC C UUCGACA		UGUCGAA CUGAUGAG X CGAA ICUGGGC	
3823	CCCAGCC U UCGACAA		UUGUCGA CUGAUGAG X CGAA IGCUGGG	
3829	CUUCGAC A ACCUCUA		UAGAGGU CUGAUGAG X CGAA IUCGAAG	
3832	CGACAAC C UCUAUUA		UAAUAGA CUGAUGAG X CGAA IUUGUCG	
3833	GACAACC U CUAUUA		GUAAUAG CUGAUGAG X CGAA IGUUGUC	
3835	CAACCUC U AUUACUG		CAGUAAU CUGAUGAG X CGAA IAGGUUG	
3841	CUAUUAC U GGGACCA		UGGUCCC CUGAUGAG X CGAA IUAAUAG	
3847	CUGGGAC C AGGACCC		GGGUCCU CUGAUGAG X CGAA IUCCCAG	
3848	UGGGACC A GGACCCA		UGGGUCC CUGAUGAG X CGAA IGUCCCA	
3853	CCAGGAC C CACCAGA		UCUGGUG CUGAUGAG X CGAA IUCCUGG	
3854	CAGGACC C ACCAGAG		CUCUGGU CUGAUGAG X CGAA IGUCCUG	
3855	AGGACCC A CCAGAGC		GCUCUGG CUGAUGAG X CGAA IGGUCCU	
3857	GACCCAC C AGAGCGG		CCGCUCU CUGAUGAG X CGAA IUGGGUC	
3858	ACCCACC A GAGCGGG		CCCGCUC CUGAUGAG X CGAA IGUGGGU	
3870	GGGGGGC U CCACCCA		UGGGUGG CUGAUGAG X CGAA ICCCCCC	

Table 34

3872	GGGGCUC C ACCCAGC		GCUGGGU CUGAUGAG X CGAA IAGCCCC	
3873	GGGCUCC A CCCAGCA		UGCUGGG CUGAUGAG X CGAA IGAGCCC	
3875	GCUCCAC C CAGCACC		GGUGCUG CUGAUGAG X CGAA IUGGAGC	
3876	CUCCACC C AGCACCU		AGGUGCU CUGAUGAG X CGAA IGUGGAG	
3877	UCCACCC A GCACCUU		AAGGUGC CUGAUGAG X CGAA IGGUGGA	
3880	ACCCAGC A CCUUCAA		UUGAAGG CUGAUGAG X CGAA ICUGGGU	
3882	CCAGCAC C UUCAAG		CUUUGAA CUGAUGAG X CGAA IUGCUGG	
3883	CAGCACC U UCAAAGG		CCUUUGA CUGAUGAG X CGAA IGUGCUG	
3886	CACCUUC A AAGGGAC		GUCCCUU CUGAUGAG X CGAA IAAGGUG	
3894	AAGGGAC A CCUACGG		CCGUAGG CUGAUGAG X CGAA IUCCCUU	
3896	GGGACAC C UACGGCA		UGCCGUA CUGAUGAG X CGAA IUGUCCC	
3897	GGACACC U ACGGCAG		CUGCCGU CUGAUGAG X CGAA IGUGUCC	
3903	CUACGGC A GAGAACC		GGUUCUC CUGAUGAG X CGAA ICCGUAG	
3910	AGAGAAC C CAGAGUA		UACUCUG CUGAUGAG X CGAA IUUCUCU	
3911	GAGAACC C AGAGUAC		GUACUCU CUGAUGAG X CGAA IGUUCUC	
3912	AGAACCC A GAGUACC		GGUACUC CUGAUGAG X CGAA IGGUUCU	
3919	AGAGUAC C UGGGUCU		AGACCCA CUGAUGAG X CGAA IUACUCU	
3920	GAGUACC U GGGUCUG		CAGACCC CUGAUGAG X CGAA IGUACUC	
3926	CUGGGUC U GGACGUG		CACGUCC CUGAUGAG X CGAA IACCCAG	
3935	GACGUGC C AGUGUGA		UCACACU CUGAUGAG X CGAA ICACGUC	
3936	ACGUGCC A GUGUGAA		UUCACAC CUGAUGAG X CGAA IGCACGU	
3945	UGUGAAC C AGAAGGC		GCCUUCU CUGAUGAG X CGAA IUUCACA	
3946	GUGAACC A GAAGGCC		GGCCUUC CUGAUGAG X CGAA IGUUCAC	
3953	AGAAGGC C AAGUCCG		CGGACU CUGAUGAG X CGAA ICCUUCU	
3954	GAAGGCC A AGUCCGC		GCGGACU CUGAUGAG X CGAA IGCCUUC	
3959	CCAAGUC C GCAGAAG		CUUCUGC CUGAUGAG X CGAA IACUUGG	
3962	AGUCCGC A GAAGCCC		GGGCUUC CUGAUGAG X CGAA ICGGACU	
3968	CAGAAGC C CUGAUGU		ACAUCAG CUGAUGAG X CGAA ICUUCUG	
3969	AGAAGCC C UGAUGUG		CACAUCA CUGAUGAG X CGAA IGCUUCU	
3970	GAAGCCC U GAUGUGU		ACACAUC CUGAUGAG X CGAA IGGCUUC	
3979	AUGUGUC C UCAGGGA		UCCUGA CUGAUGAG X CGAA IACACAU	
3980	UGUGUCC U CAGGGAG		CUCCUG CUGAUGAG X CGAA IGACACA	
3982	UGUCCUC A GGGAGCA		UGCUCU CUGAUGAG X CGAA IAGGACA	
3989	AGGGAGC A GGAAGG		CCUUCU CUGAUGAG X CGAA ICUCCU	
3998	GGAAGGC C UGACUUC		GAAGUCA CUGAUGAG X CGAA ICCUUC	
3999	GAAGGCC U GACUUCU		AGAAGUC CUGAUGAG X CGAA IGCCUUC	
4003	GCCUGAC U UCUGCUG		CAGCAGA CUGAUGAG X CGAA IUCAGGC	
4006	UGACUUC U GCUGGCA		UGCCAGC CUGAUGAG X CGAA IAAGUCA	
4009	CUUCUGC U GGCAUCA		UGAUGCC CUGAUGAG X CGAA ICAGAAG	
4013	UGCUGGC A UCAAGAG		CUCUUGA CUGAUGAG X CGAA ICCAGCA	
4016	UGGCAUC A AGAGGUG		CACCUCU CUGAUGAG X CGAA IAUGCCA	
4031	GGAGGGC C CUCCGAC		GUCGGAG CUGAUGAG X CGAA ICCCUCC	
4032	GAGGGCC C UCCGACC		GGUCGGA CUGAUGAG X CGAA IGCCUUC	
4033	AGGGCCC U CCGACCA		UGGUCGG CUGAUGAG X CGAA IGGCCCU	
4035	GGCCUUC C GACCACU		AGUGGUC CUGAUGAG X CGAA IAGGGCC	
4039	CUCCGAC C ACUCCA		UGGAAGU CUGAUGAG X CGAA IUCCGAG	
4040	UCCGACC A CUCCAG		CUGGAAG CUGAUGAG X CGAA IGUCGGA	

Table 34

4042	CGACCAC U UCCAGGG		CCCUGGA CUGAUGAG X CGAA IUGGUCCG	
4045	CCACUUC C AGGGGAA		UUCCCCU CUGAUGAG X CGAA IAAGUGG	
4046	CACUUC C A GGGGAAC		GUUCCCC CUGAUGAG X CGAA IGAAGUG	
4054	GGGGAAC C UGCCAUG		CAUGGCA CUGAUGAG X CGAA IUUCCCC	
4055	GGGAACC U GCCAUGC		GCAUGGC CUGAUGAG X CGAA IGUUCCC	
4058	AACCUGC C AUGCCAG		CUGGCAU CUGAUGAG X CGAA ICAGGUU	
4059	ACCUGCC A UGCCAGG		CCUGGCA CUGAUGAG X CGAA IGCAGGU	
4063	GCCAUGC C AGGAACC		GGUUCU CUGAUGAG X CGAA ICAUGGC	
4064	CCAUGCC A GGAACCU		AGGUUCC CUGAUGAG X CGAA IGCAUGG	
4070	CAGGAAC C UGUCCUA		UAGGACA CUGAUGAG X CGAA IUUCCUG	
4071	AGGAACC U GUCCUAA		UUAGGAC CUGAUGAG X CGAA IGUUCCU	
4075	ACCUGUC C UAAGGAA		UUCUUA CUGAUGAG X CGAA IACAGGU	
4076	CCUGUCC U AAGGAAC		GUUCCU CUGAUGAG X CGAA IGACAGG	
4084	AAGGAAC C UUCUUC		GAAGGAA CUGAUGAG X CGAA IUUCCU	
4085	AGGAACC U UCCUUC		GGAAGGA CUGAUGAG X CGAA IGUUCU	
4088	AACCUUC C UUCUGC		GCAGGAA CUGAUGAG X CGAA IAAGGUU	
4089	ACCUUC U UCCUGU		AGCAGGA CUGAUGAG X CGAA IGAAGGU	
4092	UUCUUC C UGCUUGA		UCAAGCA CUGAUGAG X CGAA IAAGGAA	
4093	UCCUUC U GCUUGAG		CUCAAGC CUGAUGAG X CGAA IGAAGGA	
4096	UUCUGC U UGAGUUC		GAACUCA CUGAUGAG X CGAA ICAGGAA	
4104	UGAGUUC C CAGAUGG		CCAUCUG CUGAUGAG X CGAA IAACUCA	
4105	GAGUUC C AGAUGGC		GCCAUCU CUGAUGAG X CGAA IGAACUC	
4106	AGUUC C A GAUGGCU		AGCCAUC CUGAUGAG X CGAA IGGAACU	
4113	AGAUGGC U GGAAGGG		CCUUC CUGAUGAG X CGAA ICCAUCU	
4124	AGGGGUC C AGCCUCG		CGAGGCU CUGAUGAG X CGAA IACCCU	
4125	GGGGUCC A GCCUCGU		ACGAGGC CUGAUGAG X CGAA IGACCCC	
4128	GUCCAGC C UCGUUGG		CCAACGA CUGAUGAG X CGAA ICUGGAC	
4129	UCCAGCC U CGUUGGA		UCCAACG CUGAUGAG X CGAA IGCUGGA	
4145	GAGGAAC A GCACUGG		CCAGUGC CUGAUGAG X CGAA IUUCCUC	
4148	GAACAGC A CUGGGGA		UCCCCAG CUGAUGAG X CGAA ICUGUUC	
4150	ACAGCAC U GGGGAGU		ACUCCCC CUGAUGAG X CGAA IUGCUGU	
4159	GGGAGUC U UUGUGGA		UCCACAA CUGAUGAG X CGAA IACUCCC	
4170	UGGAUUC U GAGGCC		GGGCCUC CUGAUGAG X CGAA IAAUCCA	
4176	CUGAGGC C CUGCCCA		UGGGCAG CUGAUGAG X CGAA ICCUCAG	
4177	UGAGGCC C UGCCCAA		UUGGGCA CUGAUGAG X CGAA IGCCUCA	
4178	GAGGCC U GCCCAAU		AUUGGGC CUGAUGAG X CGAA IGGCCUC	
4181	GCCCUGC C CAUAGAG		CUCAUUG CUGAUGAG X CGAA ICAGGGC	
4182	CCCUGCC C AAUGAGA		UCUCAU CUGAUGAG X CGAA IGCAGGG	
4183	CCUGCCC A AUGAGAC		GUCUCAU CUGAUGAG X CGAA IGGCAGG	
4191	AUGAGAC U CUAGGGU		ACCCUAG CUGAUGAG X CGAA IUCUCAU	
4193	GAGACUC U AGGGUCC		GGACCCU CUGAUGAG X CGAA IAGUCUC	
4200	UAGGGUC C AGUGGAU		AUCCACU CUGAUGAG X CGAA IACCCUA	
4201	AGGGUCC A GUGGAUG		CAUCCAC CUGAUGAG X CGAA IGACCCU	
4210	UGGAUGC C ACAGCCC		GGGCGU CUGAUGAG X CGAA ICAUCCA	
4211	GGAUGCC A CAGCCCA		UGGGCUG CUGAUGAG X CGAA IGCAUCC	
4213	AUGCCAC A GCCCAGC		GCUGGGC CUGAUGAG X CGAA IUGGCAU	
4216	CCACAGC C CAGCUUG		CAAGCUG CUGAUGAG X CGAA ICUGUGG	

Table 34

4217	CACAGCC C AGCUUGG		CCAAGCU CUGAUGAG X CGAA IGCUGUG	
4218	ACAGCCC A GCUUGGC		GCCAAGC CUGAUGAG X CGAA IGGCUGU	
4221	GCCCAGC U UGGCCCU		AGGGCCA CUGAUGAG X CGAA ICUGGGC	
4226	GCUUGGC C CUUCCU		AGGAAAG CUGAUGAG X CGAA ICCAAGC	
4227	CUUGGCC C UUCCUU		AAGGAAA CUGAUGAG X CGAA IGCCAAG	
4228	UUGGCCC U UUCUUUC		GAAGGAA CUGAUGAG X CGAA IGGCCAA	
4232	CCCUUUC C UUCAGAA		UCUGGAA CUGAUGAG X CGAA IAAAGGG	
4233	CCUUUCC U UCCAGAU		AUCUGGA CUGAUGAG X CGAA IGAAAGG	
4236	UUCCUUC C AGAUCCU		AGGAUCU CUGAUGAG X CGAA IAAGGAA	
4237	UCCUUC C GAUCCUG		CAGGAUC CUGAUGAG X CGAA IGAAGGA	
4242	CCAGAUC C UGGGUAC		GUACCCA CUGAUGAG X CGAA IAUCUGG	
4243	CAGAUC U GGGUACU		AGUACCC CUGAUGAG X CGAA IGAUCUG	
4250	UGGGUAC U GAAAGCC		GGCUUUC CUGAUGAG X CGAA IUACCCA	
4257	UGAAAGC C UUAGGGA		UCCCUAA CUGAUGAG X CGAA ICUUUCA	
4258	GAAAGCC U UAGGGAA		UUCCCUA CUGAUGAG X CGAA IGCUUUC	
4268	GGGAAGC U GGCCUGA		UCAGGCC CUGAUGAG X CGAA ICUUCCC	
4272	AGCUGGC C UGAGAGG		CCUCUCA CUGAUGAG X CGAA ICCAGCU	
4273	GCUGGCC U GAGAGGG		CCCUCUC CUGAUGAG X CGAA IGCCAGC	
4289	AAGCGGC C CUAAGGG		CCCUUAG CUGAUGAG X CGAA ICCGCUU	
4290	AGCGGCC C UAAGGGA		UCCCUUA CUGAUGAG X CGAA IGCCGCU	
4291	GCGGCCC U AAGGGAG		CUCCCUU CUGAUGAG X CGAA IGGCCGC	
4303	GAGUGUC U AAGAACA		UGUUCUU CUGAUGAG X CGAA IACACUC	
4310	UAAGAAC A AAAGCGA		UCGCUUU CUGAUGAG X CGAA IUUCUUA	
4319	AAGCGAC C CAUUCAG		CUGAAUG CUGAUGAG X CGAA IUCGCUU	
4320	AGCGACC C AUUCAGA		UCUGAAU CUGAUGAG X CGAA IGUCGCU	
4321	GCGACCC A UUCAGAG		CUCUGAA CUGAUGAG X CGAA IGGUCGC	
4325	CCCAUUC A GAGACUG		CAGUCUC CUGAUGAG X CGAA IAAUGGG	
4331	CAGAGAC U GUCCUUG		CAGGGAC CUGAUGAG X CGAA IUCUCUG	
4335	GACUGUC C CUGAAAC		GUUUCAG CUGAUGAG X CGAA IACAGUC	
4336	ACUGUCC C UGAAACC		GGUUCUA CUGAUGAG X CGAA IGACAGU	
4337	CUGUCCC U GAAACCU		AGGUUUC CUGAUGAG X CGAA IGGACAG	
4343	CUGAAAC C UAGUACU		AGUACUA CUGAUGAG X CGAA IUUUCAG	
4344	UGAAACC U AGUACUG		CAGUACU CUGAUGAG X CGAA IGUUCUA	
4350	CUAGUAC U GCCCCC		GGGGGGC CUGAUGAG X CGAA IUACUAG	
4353	GUACUGC C CCCAUG		CAUGGGG CUGAUGAG X CGAA ICAGUAC	
4354	UACUGCC C CCCAUGA		UCAUGGG CUGAUGAG X CGAA IGCAGUA	
4355	ACUGCCC C CCAUGAG		CUCAUGG CUGAUGAG X CGAA IGGCAGU	
4356	CUGCCCC C CAUGAGG		CCUCAUG CUGAUGAG X CGAA IGGGCAG	
4357	UGCCCCC C AUGAGGA		UCCUCAU CUGAUGAG X CGAA IGGGGCA	
4358	GCCCCC A UGAGGAA		UUCCUCA CUGAUGAG X CGAA IGGGGGC	
4371	AAGGAAC A GCAAUGG		CCAUUGC CUGAUGAG X CGAA IUUCCUU	
4374	GAACAGC A AUGGUGU		ACACCAU CUGAUGAG X CGAA ICUGUUC	
4383	UGGUGUC A GUAUCCA		UGGAUAC CUGAUGAG X CGAA IACACCA	
4389	CAGUAC C AGGCUUU		AAAGCCU CUGAUGAG X CGAA IAUACUG	
4390	AGUAUCC A GGCUUUG		CAAAGCC CUGAUGAG X CGAA IGAUACU	
4394	UCCAGGC U UUGUACA		UGUACAA CUGAUGAG X CGAA ICCUGGA	
4401	UUUGUAC A GAGUGCU		AGCACUC CUGAUGAG X CGAA IUACAAA	

Table 34

4408	AGAGUGC U UUUCUGU		ACAGAAA CUGAUGAG X CGAA ICACUCU	
4413	GCUUUUC U GUUUAGU		ACUAAAC CUGAUGAG X CGAA IAAAAGC	
4427	UUUUUAC U UUUUUUG		CAAAAAA CUGAUGAG X CGAA IUAAAAA	
4464	UAAAGAC C CAGGGGG		CCCCCUG CUGAUGAG X CGAA IUCUUUA	
4465	AAAGACC C AGGGGGA		UCCCCCU CUGAUGAG X CGAA IGUCUUU	
4466	AAGACCC A GGGGGAG		CUCCCCC CUGAUGAG X CGAA IGGUCUU	

Seq Accession No. = HSERB2R (Human c-erb-B-2 mRNA; 4473 bp)

Core Sequence = CUGAUGAG X CGAA (X = GCCGAAAGGC or other stem II)

Table 35

Table 35: HBV Strains and Accession numbers

Accession Number	Name
AF100308.1	AF100308 Hepatitis B virus strain 2-18, complete
AB026815.1	AB026815 Hepatitis B virus DNA, complete genome,
AB033559.1	AB033559 Hepatitis B virus DNA, complete genome,
AB033558.1	AB033558 Hepatitis B virus DNA, complete genome,
AB033557.1	AB033557 Hepatitis B virus DNA, complete genome,
AB033556.1	AB033556 Hepatitis B virus DNA, complete genome,
AB033555.1	AB033555 Hepatitis B virus DNA, complete genome,
AB033554.1	AB033554 Hepatitis B virus DNA, complete genome,
AB033553.1	AB033553 Hepatitis B virus DNA, complete genome,
AB033552.1	AB033552 Hepatitis B virus DNA, complete genome,
AB033551.1	AB033551 Hepatitis B virus DNA, complete genome,
AB033550.1	AB033550 Hepatitis B virus DNA, complete genome
AF143308.1	AF143308 Hepatitis B virus clone WB1254, complete
AF143307.1	AF143307 Hepatitis B virus clone RM518, complete
AF143306.1	AF143306 Hepatitis B virus clone RM517, complete
AF143305.1	AF143305 Hepatitis B virus clone RM501, complete
AF143304.1	AF143304 Hepatitis B virus clone HD319, complete
AF143303.1	AF143303 Hepatitis B virus clone HD1406, complete
AF143302.1	AF143302 Hepatitis B virus clone HD1402, complete
AF143301.1	AF143301 Hepatitis B virus clone BW1903, complete
AF143300.1	AF143300 Hepatitis B virus clone 7832-G4, complete
AF143299.1	AF143299 Hepatitis B virus clone 7744-G9, complete
AF143298.1	AF143298 Hepatitis B virus clone 7720-G8, complete
AB026814.1	AB026814 Hepatitis B virus DNA, complete genome,
AB026813.1	AB026813 Hepatitis B virus DNA, complete genome,
AB026812.1	AB026812 Hepatitis B virus DNA, complete genome,
AB026811.1	AB026811 Hepatitis B virus DNA, complete genome,
AJ131956.1	HBV131956 Hepatitis B virus complete genome,
AF151735.1	AF151735 Hepatitis B virus, complete genome
AF090842.1	AF090842 Hepatitis B virus strain G5.27295, complete
AF090841.1	AF090841 Hepatitis B virus strain G4.27241, complete
AF090840.1	AF090840 Hepatitis B virus strain G3.27270, complete
AF090839.1	AF090839 Hepatitis B virus strain G2.27246, complete
AF090838.1	AF090838 Hepatitis B virus strain P1.27239, complete
Y18858.1	HBV18858 Hepatitis B virus complete genome, isolate
Y18857.1	HBV18857 Hepatitis B virus complete genome, isolate
D12980.1	HPBCG Hepatitis B virus subtype adr(SRADR) DNA,
Y18856.1	HBV18856 Hepatitis B virus complete genome, isolate
Y18855.1	HBV18855 Hepatitis B virus complete genome, isolate
AJ131133.1	HBV131133 Hepatitis B virus, complete genome, strain
X80925.1	HBVP6PCXX Hepatitis B virus (patient 6) complete
X80926.1	HBVP5PCXX Hepatitis B virus (patient 5) complete
X80924.1	HBVP4PCXX Hepatitis B virus (patient 4) complete
AF100309.1	Hepatitis B virus strain 56, complete genome

Table 35

AF068756.1	AF068756 Hepatitis B virus, complete genome
AF043593.1	AF043593 Hepatitis B virus isolate 6/89, complete
Y07587.1	HBVAYWGEN Hepatitis B virus, complete genome
D28880.1	D28880 Hepatitis B virus DNA, complete genome, strain
X98076.1	HBVDEFVP3 Hepatitis B virus complete genome with
X98075.1	HBVDEFVP2 Hepatitis B virus complete genome with
X98074.1	HBVDEFVP1 Hepatitis B virus complete genome with
X98077.1	HBVCGWITY Hepatitis B virus complete genome, wild type
X98072.1	HBVCGINSC Hepatitis B virus complete genome with
X98073.1	HBVCGINCX Hepatitis B virus complete genome with
U95551.1	U95551 Hepatitis B virus subtype ayw, complete genome
D23684.1	HPBC6T588 Hepatitis B virus (C6-TKB588) complete genome
D23683.1	HPBC5HKO2 Hepatitis B virus (C5-HBVKO2) complete genome
D23682.1	HPBB5HKO1 Hepatitis B virus (B5-HBVKO1) complete genome
D23681.1	HPBC4HST2 Hepatitis B virus (C4-HBVST2) complete genome
D23680.1	HPBB4HST1 Hepatitis B virus (B4-HBVST1) complete genome
D00331.1	HPBADW3 Hepatitis B virus genome, complete genome
D00330.1	HPBADW2 Hepatitis B virus genome, complete genome
D50489.1	HPBA11A Hepatitis B virus DNA, complete genome
D23679.1	HPBA3HMS2 Hepatitis B virus (A3-HBVMS2) complete genome
D23678.1	HPBA2HYS2 Hepatitis B virus (A2-HBVYS2) complete genome
D23677.1	HPBA1HKK2 Hepatitis B virus (A1-HBVKK2) complete genome
D16665.1	HPBADRM Hepatitis B virus DNA, complete genome
D00329.1	HPBADW1 Hepatitis B virus (HBV) genome, complete genome
X97851.1	HBVP6CSX Hepatitis B virus (patient 6) complete genome
X97850.1	HBVP4CSX Hepatitis B virus (patient 4) complete genome
X97849.1	HBVP3CSX Hepatitis B virus (patient 3) complete genome
X97848.1	HBVP2CSX Hepatitis B virus (patient 2) complete genome
X51970.1	HVHEPB Hepatitis B virus (HBV 991) complete genome
M38636.1	HPBCGADR Hepatitis B virus, subtype adr, complete genome
X59795.1	HBVAYWMC G Hepatitis B virus (ayw subtype mutant)
M38454.1	HPBADR1CG Hepatitis B virus, complete genome
M32138.1	HPBHBVAA Hepatitis B virus variant HBV-alpha1, complete
J02203.1	HPBAYW Human hepatitis B virus (subtype ayw), complete
M12906.1	HPBADRA Hepatitis B virus subtype adr, complete genome
M54923.1	HPBADWZ Hepatitis B virus (subtype adw), complete genome
L27106.1	HPBMUT Hepatitis B virus mutant complete genome

Table 36

Table 36: HBV Substrate Sequence

NT Position*	Substrate	Seq ID
82	CUAUCGUGCCCUUCUUAUC	1
101	CUACCGUCCGGCC	2
159	CUUCUCAUCU	3
184	CUUCCCUACAC	4
269	GACUCUCAGAAUGUCAACGAC	5
381	CUGUAGGCAUAAAUGGUCUG	6
401	GUUCACCAGCACCAUGCAACUUUU	7
424	UUUCACGUCUGCCUAAUCAUC	8
524	AUUUGGAGCUUC	9
562	CUGACUUCUUCCUUCUAUUC	10
649	CUCACCAUACCGCACUCA	11
667	GGCAAGCUAUUCUGUG	12
717	GGAAGUAAUUUGGAAGAC	13
758	CAGCUAUGUCA AUGUUA	14
783	CUAAAUCGGCCUAAAUCAGAC	15
812	CAUJUCCUGUCACU UUUGGAAGAG	16
887	UCCUGCUUACAGAC	17
922	CAACACUUCGGAAACUACUGUUGUAG	18
989	CUCGCCUCGCAGACGAAGGUCUC	19
1009	CAAUCGCCGCGUCGCAGAAG	20
1031	AUCUCAAUCUCGGGAAUCUCAA	21
1052	AUGJUAGUAUCCUUGGACUC	22
1072	CAUAAGGUGGGAAACUJUACUG	23
1109	CUGUACCUAUUCUUUAAAUCC	24
1127	CUGAGUGGCAAACUCCC	25
1271	CCAAUAUCUGCCCUUGGACAA	26
1297	AUUAACCAUAUUAUCCUGAACA	27
1319	AUGCAGUAAUCAJUACUCAAACUA	28
1340	AAACUAGGCAUUA	29
1370	AGGCGGGCAUUCUAUAUAAGAGAG	30
1393	GAAACUACGCGCAGCGCCUCAUUUUGU	31
1412	CAUUUUGUGGGUCACCAUA	32
1441	CAAGAGCUACAGCAUGGG	33

LOCUS HPBADR1CG 3221 bp DNA circular VRL
 06-MAR-1995
 DEFINITION Hepatitis B virus , complete genome.
 ACCESSION M38454

*The nucleotide number referred to in that table is the position of the 5' end of the oligo in this sequence.

Table 37

Table 37: Human HBV Hammerhead Ribozyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
13	CCACCACT T TCCACCAA	34	UUGGUGGA CUGAUGAG X CGAA AGUGGUGG	2543
14	CACCACTT T CCACCAAA	35	UUUGGUGG CUGAUGAG X CGAA AAGUGGUG	2544
15	ACCACTTT C CACCAAAAC	36	GUUUGGUG CUGAUGAG X CGAA AAAGUGGU	2545
25	ACCAAAC T TCAAGAT	37	AUCUUGAA CUGAUGAG X CGAA AGUUGGU	2546
27	CAAACCT T CAAGATCC	38	GGAUCUUG CUGAUGAG X CGAA AGAGUUUG	2547
28	AAACTCTT C AAGATCCC	39	GGAUCUU CUGAUGAG X CGAA AAGAGUUU	2548
34	TTCAAGAT C CCAGAGTC	40	GACUCUGG CUGAUGAG X CGAA AUCUUGAA	2549
42	CCCAGAGT C AGGGCCCT	41	AGGGCCCU CUGAUGAG X CGAA ACUCUGGG	2550
53	GGCCCTGT A CTTTCCTG	42	CAGGAAAG CUGAUGAG X CGAA ACAGGGCC	2551
56	CCTGTACT T TCCTGCTG	43	CAGCAGGA CUGAUGAG X CGAA AGUACAGG	2552
57	CTGTACTT C CCTGCTGG	44	CCAGCAGG CUGAUGAG X CGAA AAGUACAG	2553
58	TGTACTTT C CTGCTGGT	45	ACCAGCAG CUGAUGAG X CGAA AAAGUACA	2554
71	TGGTGGCT C CAGTTCAG	46	CUGAACUG CUGAUGAG X CGAA AGCCACCA	2555
76	GCTCCAGT T CAGGAACA	47	UGUUCUG CUGAUGAG X CGAA ACUGGAGC	2556
77	CTCCAGT C AGGAACAG	48	CUGUUCU CUGAUGAG X CGAA AACUGGAG	2557
97	GCCCTGCT C AGAATACT	49	AGUAUUCU CUGAUGAG X CGAA AGCAGGGC	2558
103	CTCAGAAT A CTGTCTCT	50	AGAGACAG CUGAUGAG X CGAA AUUCUGAG	2559
108	AATACTGT C TCTGCCAT	51	AUGGCAGA CUGAUGAG X CGAA ACAGUAUU	2560
110	TACTGTCT C TGCCATAT	52	AUAUGGCA CUGAUGAG X CGAA AGACAGUA	2561
117	TCTGCCAT A TCGTCAAT	53	AUUGACGA CUGAUGAG X CGAA AUGGCAGA	2562
119	TGCCATAT C GTCAATCT	54	AGAUUGAC CUGAUGAG X CGAA AUAUGGCA	2563
122	CATATCGT C AATCTTAT	55	AUAAGAUU CUGAUGAG X CGAA ACGAUUUG	2564
126	TCGTCAAT C TTATCGAA	56	UUCGAUAA CUGAUGAG X CGAA AUUGACGA	2565
128	GTCAATCT T ATCGAAGA	57	UCUUCGAU CUGAUGAG X CGAA AGAUUGAC	2566
129	TCAATCTT A TCGAAGAC	58	GUCUUCGA CUGAUGAG X CGAA AAGAUUGA	2567
131	AATCTTAT C GAAGACTG	59	CAGUCUUC CUGAUGAG X CGAA AUAAGAUU	2568
150	GACCCTGT A CCGAACAT	60	AUGUUCGG CUGAUGAG X CGAA ACAGGGUC	2569
168	GAGAACAT C GCATCAGG	61	CCUGAUGC CUGAUGAG X CGAA AUGUUCUC	2570
173	CATCGCAT C AGGACTCC	62	GGAGUCCU CUGAUGAG X CGAA AUGCGAUG	2571
180	TCAGGACT C CTAGGACC	63	GGUCCUAG CUGAUGAG X CGAA AGUCCUGA	2572
183	GGACTCCT A GGACCCCT	64	AGGGGUCC CUGAUGAG X CGAA AGGAGUCC	2573
195	CCCCTGCT C GTGTTACA	65	UGUAACAC CUGAUGAG X CGAA AGCAGGGG	2574
200	GCTCGTGT T ACAGGCGG	66	CCGCCUGU CUGAUGAG X CGAA ACACGAGC	2575
201	CTCGTGTT A CAGGCGGG	67	CCGCCUG CUGAUGAG X CGAA AACACGAG	2576
212	GGCGGGGT T TTTCTTGT	68	ACAAGAAA CUGAUGAG X CGAA ACCCCGCC	2577
213	GCGGGGTT T TTCTTGT	69	AACAAGAA CUGAUGAG X CGAA AACCCCGC	2578
214	CGGGGTTT T TCTTGTG	70	CAACAAGA CUGAUGAG X CGAA AAACCCCG	2579
215	GGGGTTTT T CTTGTTGA	71	UCAACAAG CUGAUGAG X CGAA AAAACCCC	2580
216	GGGTTTTT C TTGTTGAC	72	GUCAACAA CUGAUGAG X CGAA AAAAACCC	2581
218	GTTTTTCT T GTTGACAA	73	UUGUCAAC CUGAUGAG X CGAA AGAAAAAC	2582
221	TTCTTGT T GACAAAAA	74	UUUUUGUC CUGAUGAG X CGAA ACAAGAAA	2583
231	ACAAAAAT C CTCACAAT	75	AUUGUGAG CUGAUGAG X CGAA AUUUUUGU	2584
234	AAAAATCCT C ACAATACC	76	GGUAUUGU CUGAUGAG X CGAA AGGAUUUU	2585
240	CTCACAAT A CCACAGAG	77	CUCUGUGG CUGAUGAG X CGAA AUUGUGAG	2586
250	CACAGAGT C TAGACTCG	78	CGAGUCUA CUGAUGAG X CGAA ACUCUGUG	2587
252	CAGAGTCT A GACTCGTG	79	CACGAGUC CUGAUGAG X CGAA AGACUCUG	2588
257	TCTAGACT C GTGGTGGA	80	UCCACCAC CUGAUGAG X CGAA AGUCUAGA	2589
268	GGTGGACT T CTCTCAAT	81	AUUGAGAG CUGAUGAG X CGAA AGUCCACC	2590

Table 37

269	GTGGACTT C TCTCAATT	82	AAUUGAGA CUGAUGAG X CGAA AAGUCCAC	2591
271	GGACTTCT C TCAATTTT	83	AAAAUUGA CUGAUGAG X CGAA AGAAGUCC	2592
273	ACTTCTCT C AATTTTCT	84	AGAAAAJU CUGAUGAG X CGAA AGAGAAGU	2593
277	CTCTCAAT T TTCTAGGG	85	CCCUAGAA CUGAUGAG X CGAA AUUGAGAG	2594
278	TCTCAATT T TCTAGGGG	86	CCCCUAGA CUGAUGAG X CGAA AAUUGAGA	2595
279	CTCAATTT T CTAGGGGG	87	CCCCCUAG CUGAUGAG X CGAA AAAUUGAG	2596
280	TCAATTTT C TAGGGGGA	88	UCCCCCUA CUGAUGAG X CGAA AAAAUUGA	2597
282	AATTTTCT A GGGGGAAC	89	GUCCCCC CUGAUGAG X CGAA AGAAAAUU	2598
301	CCGTGTGT C TTGGCCAA	90	UUGGCCAA CUGAUGAG X CGAA ACACACGG	2599
303	GTGTGTCT T GGCCAAAA	91	UUUUGGCC CUGAUGAG X CGAA AGACACAC	2600
313	GCCAAAAT T CGCAGTCC	92	GGACUGCG CUGAUGAG X CGAA AUUUUGGC	2601
314	CCAAAATT C GCAGTCCC	93	GGGACUGC CUGAUGAG X CGAA AAUUUUGG	2602
320	TTCGCACT C CCAAATCT	94	AGAUUUGG CUGAUGAG X CGAA ACUGCGAA	2603
327	TCCCAAAT C TCCAGTCA	95	UGACUGGA CUGAUGAG X CGAA AUUUGGGA	2604
329	CCAAATCT C CAGTCACT	96	AGUGACUG CUGAUGAG X CGAA AGAUUUGG	2605
334	TCTCCAGT C ACTCACCA	97	UGGUGAGU CUGAUGAG X CGAA ACUGGAGA	2606
338	CAGTCACT C ACCAACCT	98	AGGUUGGU CUGAUGAG X CGAA AGUGACUG	2607
349	CAACCTGT T GTCCTCCA	99	UGGAGGAC CUGAUGAG X CGAA ACAGGUUG	2608
352	CCTGTTGT C CTCCAATT	100	AAUUGGAG CUGAUGAG X CGAA ACAACAGG	2609
355	GTTGTCTT C CAATTGT	101	ACAAAUUG CUGAUGAG X CGAA AGGACAAC	2610
360	CCTCCAAT T TGTCTGG	102	CCAGGACA CUGAUGAG X CGAA AUUGGAGG	2611
361	CTCCAATT T GTCCTGGT	103	ACCAGGAC CUGAUGAG X CGAA AAUUGGAG	2612
364	CAATTGT C CTGTTAT	104	AUAACCAG CUGAUGAG X CGAA ACAAUUG	2613
370	GTCCTGGT T ATCGCTGG	105	CCAGCGAU CUGAUGAG X CGAA ACCAGGAC	2614
371	TCCTGGTT A TCGCTGGA	106	UCCAGCGA CUGAUGAG X CGAA AACCAGGA	2615
373	CTGTTAT C GCTGGATG	107	CAUCCAGC CUGAUGAG X CGAA AUAACCAG	2616
385	GGATGTGT C TGC GCGT	108	ACGCCCA CUGAUGAG X CGAA ACACAUCC	2617
394	TGCGGCGT T TTATCATC	109	GAUGAUAA CUGAUGAG X CGAA ACGCCCA	2618
395	GCGGCGTT T TATCATCT	110	AGAUGAUA CUGAUGAG X CGAA AACGCCGC	2619
396	CGGCGTTT T ATCATCTT	111	AAGAUGAU CUGAUGAG X CGAA AAACGCCG	2620
397	GGCGTTTT A TCATCTTC	112	GAAGAUA CUGAUGAG X CGAA AAAACGCC	2621
399	CGTTTTAT C ATCTTCCT	113	AGGAAGAU CUGAUGAG X CGAA AUAACG	2622
402	TTTATCAT C TTCCTCTG	114	CAGAGGAA CUGAUGAG X CGAA AUGAUAAA	2623
404	TATCATCT T CCTCTGCA	115	UGCAGAGG CUGAUGAG X CGAA AGAUGAUA	2624
405	ATCATCTT C CTCTGCAT	116	AUGCAGAG CUGAUGAG X CGAA AAGAUGAU	2625
408	ATCTTCCT C TGCATCCT	117	AGGAUGCA CUGAUGAG X CGAA AGGAAGAU	2626
414	CTCTGCAT C CTGCTGCT	118	AGCAGCAG CUGAUGAG X CGAA AUGCAGAG	2627
423	CTGCTGCT A TGCCTCAT	119	AUGAGGCA CUGAUGAG X CGAA AGCAGCAG	2628
429	CTATGCCT C ATCTTCTT	120	AAGAAGAU CUGAUGAG X CGAA AGGCAUAG	2629
432	TGCCTCAT C TTCTTGT	121	AACAAGAA CUGAUGAG X CGAA AUGAGGCA	2630
434	CCTCATCT T CTGTTGG	122	CCAACAAG CUGAUGAG X CGAA AGAUGAGG	2631
435	CTCATCTT C TTGTTGGT	123	ACCAACAA CUGAUGAG X CGAA AAGAUGAG	2632
437	CATCTTCT T GTTGGTTC	124	GAACCAAC CUGAUGAG X CGAA AGAAGAUG	2633
440	CTTCTTGT T GGTCTTTC	125	GAAGAACC CUGAUGAG X CGAA ACAAGAAG	2634
444	TTGTTGGT T CTTCTGGA	126	UCCAGAAG CUGAUGAG X CGAA ACCAACAA	2635
445	TGTTGGTT C TTCTGGAC	127	GUCCAGAA CUGAUGAG X CGAA AACCAACA	2636
447	TTGGTTCT T CTGACTA	128	UAGUCCAG CUGAUGAG X CGAA AGAACCAA	2637
448	TGGTTCTT C TGGACTAT	129	AUAGUCCA CUGAUGAG X CGAA AAGAACCA	2638
455	TCTGGACT A TCAAGGTA	130	UACCUUGA CUGAUGAG X CGAA AGUCCAGA	2639
457	TGGACTAT C AAGGTATG	131	CAUACCUU CUGAUGAG X CGAA AUAGUCCA	2640
463	ATCAAGGT A TGTGCCC	132	GGCAACA CUGAUGAG X CGAA ACCUUGAU	2641

Table 3/

467	AGGTATGT T GCCCGTTT	133	AAACGGGC CUGAUGAG X CGAA ACAUACCU	2642
474	TTGCCCGT T TGTCTCT	134	AGAGGACA CUGAUGAG X CGAA ACGGGCAA	2643
475	TGCCCGTT T GTCCTCTA	135	UAGAGGAC CUGAUGAG X CGAA AACGGGCA	2644
478	CCGTTTGT C CTCTAATT	136	AAUUGAG CUGAUGAG X CGAA ACAAACGG	2645
481	TTTGTCT C TAATTCCA	137	UGGAAUUA CUGAUGAG X CGAA AGGACAAA	2646
483	TGTCTCT A ATTCCAGG	138	CCUGGAAU CUGAUGAG X CGAA AGAGGACA	2647
486	CCTCTAAT T CCAGGATC	139	GAUCCUGG CUGAUGAG X CGAA AUUAGAGG	2648
487	CTCTAATT C CAGGATCA	140	UGAUCCUG CUGAUGAG X CGAA AAUUGAG	2649
494	TCCAGGAT C ATCAACAA	141	UUGUUGAU CUGAUGAG X CGAA AUCCUGGA	2650
497	AGGATCAT C AACAACCA	142	UGGUUGUU CUGAUGAG X CGAA AUGAUCCU	2651
535	GCACAACT C CTGCTCAA	143	UUGAGCAG CUGAUGAG X CGAA AGUUGUGC	2652
541	CTCCTGCT C AAGGAACC	144	GGUCCUU CUGAUGAG X CGAA AGCAGGAG	2653
551	AGGAACCT C TATGTTTC	145	GAAACAUA CUGAUGAG X CGAA AGGUCCU	2654
553	GAACCTCT A TGTTCCC	146	GGGAAACA CUGAUGAG X CGAA AGAGGUUC	2655
557	CTCTATGT T TCCTCAT	147	AUGAGGGA CUGAUGAG X CGAA ACAUAGAG	2656
558	TCTATGTT T CCCTCATG	148	CAUGAGGG CUGAUGAG X CGAA AACAUAGA	2657
559	CTATGTTT C CCTCATGT	149	ACAUGAGG CUGAUGAG X CGAA AAACUAG	2658
563	GTTTCCCT C ATGTTGCT	150	AGCAACAU CUGAUGAG X CGAA AGGGAAAC	2659
568	CCTCATGT T GCTGTACA	151	UGUACAG CUGAUGAG X CGAA ACAUGAGG	2660
574	GTTGCTGT A CAAACCT	152	AGGUUUUG CUGAUGAG X CGAA ACAGCAAC	2661
583	CAAAACCT A CGGACGGA	153	UCCGUCCG CUGAUGAG X CGAA AGGUUUUG	2662
604	GCACCTGT A TTCCATC	154	GAUGGAA CUGAUGAG X CGAA ACAGGUCC	2663
606	ACCTGTAT T CCCATCCC	155	GGGAUGGG CUGAUGAG X CGAA AUACAGGU	2664
607	CCTGTATT C CCATCCCA	156	UGGGAUGG CUGAUGAG X CGAA AAUACAGG	2665
612	ATTCCCAT C CCATCATC	157	GAUGAUGG CUGAUGAG X CGAA AUGGGAAU	2666
617	CATCCCAT C ATCTGGG	158	CCCAAGAU CUGAUGAG X CGAA AUGGGAUG	2667
620	CCCATCAT C TTGGGCTT	159	AAGCCCAA CUGAUGAG X CGAA AUGAUGGG	2668
622	CATCATCT T GGGCTTTC	160	GAAAGCCC CUGAUGAG X CGAA AGAUGAUG	2669
628	CTTGGGCT T TCGCAAAA	161	UUUUGCGA CUGAUGAG X CGAA AGCCCAAG	2670
629	TTGGGCTT T CGCAAAAT	162	AUUUUGCG CUGAUGAG X CGAA AAGCCCAA	2671
630	TGGGCTTT C GCAAAATA	163	UAUUUUGC CUGAUGAG X CGAA AAAGCCCA	2672
638	CGCAAAAT A CCTATGGG	164	CCCAUAGG CUGAUGAG X CGAA AUUUUGCG	2673
642	AAATACCT A TGGGAGTG	165	CACUCCCA CUGAUGAG X CGAA AGGUUUUU	2674
656	GTGGGCCT C AGTCCGTT	166	AACGGACU CUGAUGAG X CGAA AGGCCAC	2675
660	GCCTCAGT C CGTTCTC	167	GAGAAACG CUGAUGAG X CGAA ACUGAGGC	2676
664	CAGTCCGT T TCTCTGG	168	CCAAGAGA CUGAUGAG X CGAA ACGGACUG	2677
665	AGTCCGTT T CTCTGGC	169	GCCAAGAG CUGAUGAG X CGAA AACGGACU	2678
666	GTCCGTTT C TCTTGGCT	170	AGCCAAGA CUGAUGAG X CGAA AAACGGAC	2679
668	CCGTTTCT C TTGGCTCA	171	UGAGCCAA CUGAUGAG X CGAA AGAAACGG	2680
670	GTTTCTCT T GGCTCAGT	172	ACUGAGCC CUGAUGAG X CGAA AGAGAAAC	2681
675	TCTTGGCT C AGTTTACT	173	AGUAAACU CUGAUGAG X CGAA AGCCAAGA	2682
679	GGCTCAGT T TACTAGTG	174	CACUAGUA CUGAUGAG X CGAA ACUGAGCC	2683
680	GCTCAGTT T ACTAGTGC	175	GCACUAGU CUGAUGAG X CGAA AACUGAGC	2684
681	CTCAGTTT A CTAGTGCC	176	GGCACUAG CUGAUGAG X CGAA AACUGAG	2685
684	AGTTTACT A GTGCCATT	177	AAUGGCAC CUGAUGAG X CGAA AGUAAACU	2686
692	AGTGCCAT T TGTTCACT	178	ACUGAACA CUGAUGAG X CGAA AUGGCACU	2687
693	GTGCCATT T GTTCAGTG	179	CACUGAAC CUGAUGAG X CGAA AAUGGCAC	2688
696	CCATTTGT T CAGTGGTT	180	AACCACUG CUGAUGAG X CGAA ACAAUUGG	2689
697	CATTTGTT C AGTGGTTC	181	GAACCACU CUGAUGAG X CGAA AACAAUUG	2690
704	TCAGTGGT T CGTAGGGC	182	GCCCUACG CUGAUGAG X CGAA ACCACUGA	2691
705	CAGTGGTT C GTAGGGCT	183	AGCCCUAC CUGAUGAG X CGAA AACCACUG	2692

Table 37

708	TGGTTCGT A GGGCTTTC	184	GAAAGCCC CUGAUGAG X CGAA ACGAACCA	2693
714	GTAGGGCT T TCCCCAC	185	GUGGGGGA CUGAUGAG X CGAA AGCCCUAC	2694
715	TAGGGCTT T CCCCCACT	186	AGUGGGGG CUGAUGAG X CGAA AAGCCCUA	2695
716	AGGGCTTT C CCCCCACTG	187	CAGUGGGG CUGAUGAG X CGAA AAAGCCCU	2696
726	CCCACTGT C TGGCTTTC	188	GAAAGCCA CUGAUGAG X CGAA ACAGUGGG	2697
732	GTCTGGCT T TCAGTTAT	189	AUAACUGA CUGAUGAG X CGAA AGCCAGAC	2698
733	TCTGGCTT T CAGTTATA	190	UAUAACUG CUGAUGAG X CGAA AAGCCAGA	2699
734	CTGGCTTT C AGTTATAT	191	AUAUAACU CUGAUGAG X CGAA AAAGCCAG	2700
738	CTTTCAGT T ATATGGAT	192	AUCCAUAU CUGAUGAG X CGAA ACUGAAAG	2701
739	TTTCAGTT A TATGGATG	193	CAUCCAUA CUGAUGAG X CGAA AACUGAAA	2702
741	TCAGTTAT A TGGATGAT	194	AUCAUCCA CUGAUGAG X CGAA AUAACUGA	2703
755	GATGTGGT T TTGGGGGC	195	GCCCCCAA CUGAUGAG X CGAA ACCACAUC	2704
756	ATGTGGTT T TGGGGGCC	196	GGCCCCCA CUGAUGAG X CGAA AACCACAU	2705
757	TGTGGTTT T GGGGGCCA	197	UGGCCCCC CUGAUGAG X CGAA AAACCACA	2706
769	GGCCAAGT C TGTACAAC	198	GUUGUACA CUGAUGAG X CGAA ACUUGGCC	2707
773	AAGTCTGT A CAACATCT	199	AGAUGUUG CUGAUGAG X CGAA ACAGACUU	2708
780	TACAACAT C TTGAGTCC	200	GGACUCAA CUGAUGAG X CGAA AUGUUGUA	2709
782	CAACATCT T GAGTCCCT	201	AGGGACUC CUGAUGAG X CGAA AGAUGUUG	2710
787	TCTTGAGT C CCTTTATG	202	CAUAAAGG CUGAUGAG X CGAA ACUCAAGA	2711
791	GAGTCCCT T TATGCCGC	203	GCGGCAUA CUGAUGAG X CGAA AGGGACUC	2712
792	AGTCCCTT T ATGCCGCT	204	AGCGGCAU CUGAUGAG X CGAA AAGGGACU	2713
793	GTCCCTTT A TGCCGCTG	205	CAGCGGCA CUGAUGAG X CGAA AAAGGGAC	2714
803	GCCGCTGT T ACCAATTT	206	AAAUUGGU CUGAUGAG X CGAA ACAGCGGC	2715
804	CCGCTGTT A CCAATTTT	207	AAAAUUGG CUGAUGAG X CGAA AACAGCGG	2716
810	TTACCAAT T TTCTTTTG	208	CAAAAGAA CUGAUGAG X CGAA AUUGGUAA	2717
811	TACCAATT T TCTTTTGT	209	ACAAAAGA CUGAUGAG X CGAA AAUUGGUA	2718
812	ACCAATTT T CTTTGTCT	210	GACAAAAG CUGAUGAG X CGAA AAAUUGGU	2719
813	CCAATTTT C TTTTGTCT	211	AGACAAA A CUGAUGAG X CGAA AAAAUUGG	2720
815	AATTTTCT T TTGTCTTT	212	AAAGACAA CUGAUGAG X CGAA AGAAAAUU	2721
816	ATTTTCTT T TGTCTTTG	213	CAAAGACA CUGAUGAG X CGAA AAGAAAAU	2722
817	TTTTCTTT T GTCTTTGG	214	CCAAAGAC CUGAUGAG X CGAA AAAGAAAA	2723
820	TCTTTTGT C TTGGGTATA	215	UACCCAAA CUGAUGAG X CGAA ACAAAGA	2724
822	TTTTGTCT T TGGGTATA	216	UAUACCCA CUGAUGAG X CGAA AGACAAAA	2725
823	TTTGTCTT T GGGTATAC	217	GUUAACCC CUGAUGAG X CGAA AAGACAAA	2726
828	CTTTGGGT A TACATTTA	218	UAAAUUGA CUGAUGAG X CGAA ACCCAAAG	2727
830	TTGGGTAT A CATTTAAA	219	UTUAAAUG CUGAUGAG X CGAA AUACCCAA	2728
834	GTATACAT T TAAACCCT	220	AGGGUUUA CUGAUGAG X CGAA AUGUAUAC	2729
835	TATACATT T AAACCCTC	221	GAGGGUUU CUGAUGAG X CGAA AAUGUAUA	2730
836	ATACATTT A AACCCTCA	222	UGAGGGUU CUGAUGAG X CGAA AAAUGUAU	2731
843	TAAACCCT C ACAAACA	223	UGUUUUGU CUGAUGAG X CGAA AGGGUUUA	2732
865	ATGGGGAT A TTCCCTTA	224	UAAGGGAA CUGAUGAG X CGAA AUCCCAU	2733
867	GGGGATAT T CCCTTAAC	225	GUUAAGGG CUGAUGAG X CGAA AUAUCCCC	2734
868	GGGATATT C CCTTAAC	226	AGUUAAGG CUGAUGAG X CGAA AAUAUCCC	2735
872	TATTCCCT T AACTTCAT	227	AUGAAGUU CUGAUGAG X CGAA AGGGAAUA	2736
873	ATTCCCTT A ACTTCATG	228	CAUGAAGU CUGAUGAG X CGAA AAGGGAAU	2737
877	CCTTAAC T CATGGGAT	229	AUCCCAUG CUGAUGAG X CGAA AGUUAAGG	2738
878	CTTAACCT C ATGGGATA	230	UAUCCCAU CUGAUGAG X CGAA AAGUUAAG	2739
886	CATGGGAT A TGTAATTG	231	CAAUUACA CUGAUGAG X CGAA AUCCCAUG	2740
890	GGATATGT A ATTGGGAG	232	CUCCCAAU CUGAUGAG X CGAA ACAUAUCC	2741
893	TATGTAAT T GGGAGTTG	233	CAACUCCC CUGAUGAG X CGAA AUUACAA	2742
900	TTGGGAGT T GGGGCACA	234	UGUGCCCC CUGAUGAG X CGAA ACUCCCAA	2743

Table 3/

910	GGGCACAT T GCCACAGG	235	CCUGUGGC CUGAUGAG X CGAA AUGUGCCC	2744
924	AGGAACAT A TTGTACAA	236	UUGUACAA CUGAUGAG X CGAA AUGUCCU	2745
926	GAACATAT T GTACAAAA	237	UUUUGUAC CUGAUGAG X CGAA AUAUGUUC	2746
929	CATATTGT A CAAAAAAT	238	AUUUUUUG CUGAUGAG X CGAA ACAUAUG	2747
938	CAAAAAAT C AAAATGTG	239	CACAUUUU CUGAUGAG X CGAA AUUUUUUG	2748
948	AAATGTGT T TTAGGAAA	240	UUUCCUAA CUGAUGAG X CGAA ACACAUUU	2749
949	AATGTGTT T TAGGAAAC	241	GUUCCUA CUGAUGAG X CGAA AACACAUU	2750
950	ATGTGTTT T AGGAAACT	242	AGUUCCU CUGAUGAG X CGAA AAACACAU	2751
951	TGTGTTTT A GGAACTT	243	AAGUUCC CUGAUGAG X CGAA AAAACACA	2752
959	AGGAAACT T CCTGTAAA	244	UUUACAGG CUGAUGAG X CGAA AGUUCCU	2753
960	GGAAACTT C CTGTAAAC	245	GUUUACAG CUGAUGAG X CGAA AAGUUCC	2754
965	CTTCTGT A AACAGGCC	246	GGCCUGUU CUGAUGAG X CGAA ACAGGAAG	2755
975	ACAGGCCT A TTGATTGG	247	CCAAUCAA CUGAUGAG X CGAA AGGCCUGU	2756
977	AGGCCTAT T GATTGGAA	248	UUCCAAUC CUGAUGAG X CGAA AUAGCCU	2757
981	CTATTGAT T GGAAAGTA	249	UACUUCC CUGAUGAG X CGAA AUCAAUAG	2758
989	TGGAAGT A TGTCACG	250	CGUUGACA CUGAUGAG X CGAA ACUUCCA	2759
993	AAGTATGT C AACGAATT	251	AAUUCGU CUGAUGAG X CGAA ACAUACU	2760
1001	CAACGAAT T GTGGTCT	252	AGACCAC CUGAUGAG X CGAA AUUCGUUG	2761
1008	TTGTGGGT C TTTGGGG	253	CCCCAAA CUGAUGAG X CGAA ACCCACA	2762
1010	GTGGGTCT T TTGGGGT	254	AACCCCA CUGAUGAG X CGAA AGACCAC	2763
1011	TGGGTCTT T TGGGGTT	255	AAACCCA CUGAUGAG X CGAA AAGACCA	2764
1012	GGGTCTTT T GGGGTTG	256	CAAACCC CUGAUGAG X CGAA AAAGACC	2765
1018	TTTGGGT T TGCCGCC	257	GGGCGCA CUGAUGAG X CGAA ACCCCAA	2766
1019	TTGGGGT T GCCGCC	258	GGGCGGC CUGAUGAG X CGAA AACCCAA	2767
1029	CGCCCTT T TCACGCA	259	UUGCGUA CUGAUGAG X CGAA AGGGCGG	2768
1030	CGCCCTT T CACGCAAT	260	AUUGCGUG CUGAUGAG X CGAA AAGGGCG	2769
1031	GCCCTTT C ACGCAATG	261	CAUUGCGU CUGAUGAG X CGAA AAAGGGC	2770
1045	ATGTGGAT A TTCTGCTT	262	AAGCAGAA CUGAUGAG X CGAA AUCCACAU	2771
1047	GTGGATAT T CTGCTTAA	263	UAAAGCAG CUGAUGAG X CGAA AUAUCCAC	2772
1048	TGGATATT C TGCTTAA	264	UUAAAGCA CUGAUGAG X CGAA AAUAUCCA	2773
1053	ATTCTGCT T TAATGCCT	265	AGGCAUUA CUGAUGAG X CGAA AGCAGAAU	2774
1054	TTCTGCTT T AATGCCTT	266	AAGGCAU CUGAUGAG X CGAA AAGCAGAA	2775
1055	TCTGCTTT A ATGCCTTT	267	AAAGGCAU CUGAUGAG X CGAA AAAGCAGA	2776
1062	TAATGCCT T TATATGCA	268	UGCAUUA CUGAUGAG X CGAA AGGCAUUA	2777
1063	AATGCCTT T ATATGCAT	269	AUGCAUUA CUGAUGAG X CGAA AAGGCAU	2778
1064	ATGCCTTT A TATGCATG	270	CAUGCAUA CUGAUGAG X CGAA AAAGGCAU	2779
1066	GCCTTTAT A TGCATGCA	271	UGCAUGCA CUGAUGAG X CGAA AUAAGGC	2780
1076	GCATGCAT A CAAGCAAA	272	UUUGCUUG CUGAUGAG X CGAA AUGCAUGC	2781
1092	AACAGGCT T TTAATTCT	273	GAAAGUAA CUGAUGAG X CGAA AGCCUGU	2782
1093	ACAGGCTT T TACTTTCT	274	AGAAAGUA CUGAUGAG X CGAA AAGCCUGU	2783
1094	CAGGCTTT T ACTTTCTC	275	GAGAAAGU CUGAUGAG X CGAA AAAGCCUG	2784
1095	AGGCTTTT A CTTTCTCG	276	CGAGAAAG CUGAUGAG X CGAA AAAAGCCU	2785
1098	CTTTTACT T TCTCGCCA	277	UGGCGAGA CUGAUGAG X CGAA AGUAAAAG	2786
1099	TTTACTTT T TCTGCCAA	278	UUGGCGAG CUGAUGAG X CGAA AAGUAAA	2787
1100	TTTACTTT C TCGCCAAC	279	GUUGGCGA CUGAUGAG X CGAA AAAGUAAA	2788
1102	TACTTTCT C GCCAACTT	280	AAGUUGGC CUGAUGAG X CGAA AGAAAGUA	2789
1110	CGCCAACT T ACAAGGCC	281	GGCCUUGU CUGAUGAG X CGAA AGUUGGCG	2790
1111	GCCAACTT A CAAGGCCT	282	AGGCCUUG CUGAUGAG X CGAA AAGUUGGC	2791
1120	CAAGGCCT T TCTAAGTA	283	UACUUAGA CUGAUGAG X CGAA AGGCCUUG	2792
1121	AAGGCCTT T CTAAGTAA	284	UUACUUAG CUGAUGAG X CGAA AAGGCCU	2793
1122	AGGCCTTT C TAAGTAAA	285	UUUACUUA CUGAUGAG X CGAA AAAGGCCU	2794

Table 37

1124	GCCTTTCT A AGTAAACA	286	UGUUUACU CUGAUGAG X CGAA AGAAAGGC	2795
1128	TTCTAAGT A AACAGTAT	287	AUACUGUU CUGAUGAG X CGAA ACUUGAGAA	2796
1135	TAAACAGT A TGTGAACC	288	GGUUCACA CUGAUGAG X CGAA ACUGUUUA	2797
1145	GTGAACCT T TACCCCGT	289	ACGGGGUA CUGAUGAG X CGAA AGGUTUCAC	2798
1146	TGAACCTT T ACCCCGTT	290	AACGGGGU CUGAUGAG X CGAA AAGGUUCA	2799
1147	GAACCTTT A CCCCGTTG	291	CAACGGGG CUGAUGAG X CGAA AAAGGUUC	2800
1154	TACCCCGT T GCTCGGCA	292	UGCCGAGC CUGAUGAG X CGAA ACGGGGUA	2801
1158	CCGTTGCT C GGCAACGG	293	CCGUUGCC CUGAUGAG X CGAA AGCAACGG	2802
1173	GGCCTGGT C TATGCCAA	294	UUGGCAUA CUGAUGAG X CGAA ACCAGGCC	2803
1175	CCTGGTCT A TGCCAAGT	295	ACUUGGCA CUGAUGAG X CGAA AGACCAGG	2804
1186	CCAAGTGT T TGCTGACG	296	CGUCAGCA CUGAUGAG X CGAA ACACUUGG	2805
1187	CAAGTGT T GCTGACGC	297	GCGUCAGC CUGAUGAG X CGAA AACACUUG	2806
1209	CCACTGGT T GGGGCTTG	298	CAAGCCCC CUGAUGAG X CGAA ACCAGUGG	2807
1216	TTGGGGCT T GGCCATAG	299	CUAUGGCC CUGAUGAG X CGAA AGCCCCAA	2808
1223	TTGGCCAT A GGCCATCA	300	UGAUGGCC CUGAUGAG X CGAA AUGGCCAA	2809
1230	TAGGCCAT C AGCGCATG	301	CAUGCGCU CUGAUGAG X CGAA AUGGCCUA	2810
1249	TGGAACCT T TGTGTCTC	302	GAGACACA CUGAUGAG X CGAA AGGUUCCA	2811
1250	GGAACTT T GTGTCTCC	303	GGAGACAC CUGAUGAG X CGAA AAGGUUCC	2812
1255	CTTTGTGT C TCCTCTGC	304	GCAGAGGA CUGAUGAG X CGAA ACACAAAG	2813
1257	TTGTGTCT C CTCTGCCG	305	CGGCAGAG CUGAUGAG X CGAA AGACACAA	2814
1260	TGTCTCCT C TGCCGATC	306	GAUCGGCA CUGAUGAG X CGAA AGGAGACA	2815
1268	CTGCCGAT C CATACCGC	307	GCGGUUAG CUGAUGAG X CGAA AUCGGCAG	2816
1272	CGATCCAT A CCGCGGAA	308	UUCGCGG CUGAUGAG X CGAA AUGGAUCG	2817
1283	GCGGAAC T C TAGCCGC	309	GCGGUUAG CUGAUGAG X CGAA AGUUCGCG	2818
1286	GAACCTCT A GCCGCTTG	310	CAAGCGGC CUGAUGAG X CGAA AGGAGUUC	2819
1293	TAGCCGCT T GTTTTGCT	311	AGCAAAAC CUGAUGAG X CGAA AGCGGCUA	2820
1296	CCGCTTGT T TTGCTCGC	312	GCGAGCAA CUGAUGAG X CGAA ACAAGCGG	2821
1297	CGCTTGTT T TGCTCGCA	313	UGCGAGCA CUGAUGAG X CGAA AACAGCGG	2822
1298	GCTTGTTT T GCTCGCAG	314	CUGCGAGC CUGAUGAG X CGAA AAACAAGC	2823
1302	GTTTGCT C GCAGCAGG	315	CCUGCUGC CUGAUGAG X CGAA AGCAAAAC	2824
1312	CAGCAGT C TGGGGCAA	316	UUGCCCCA CUGAUGAG X CGAA ACCUGCUG	2825
1325	GCAAACT C ATCGGGAC	317	GUCCCCAU CUGAUGAG X CGAA AGUUUUGC	2826
1328	AAATCAT C GGGACTGA	318	UCAGUCCC CUGAUGAG X CGAA AUGAGUUU	2827
1341	CTGACAAT T CTGTCGTG	319	CACGACAG CUGAUGAG X CGAA AUUGUCAG	2828
1342	TGACAATT C TGTCGTGC	320	GCACGACA CUGAUGAG X CGAA AAUUGUCA	2829
1346	AATTCTGT C GTGCTCTC	321	GAGAGCAC CUGAUGAG X CGAA ACAGAAUU	2830
1352	GTCGTGCT C TCCGCAA	322	UUGCGGGA CUGAUGAG X CGAA AGCACGAC	2831
1354	CGTGCTCT C CCGCAAAT	323	AUUUGCGG CUGAUGAG X CGAA AGAGCACG	2832
1363	CCGCAAAT A TACATCAT	324	AUGAUGUA CUGAUGAG X CGAA AUUUGCGG	2833
1365	GCAAATAT A CATCATTT	325	AAAUGAUG CUGAUGAG X CGAA AUUUUUGC	2834
1369	ATATACAT C ATTTCCAT	326	AUGGAAAU CUGAUGAG X CGAA AUGUAUUA	2835
1372	TACATCAT T TCCATGGC	327	GCCAUGGA CUGAUGAG X CGAA AUGAUGUA	2836
1373	ACATCATT T CCATGGCT	328	AGCCAUGG CUGAUGAG X CGAA AAUGAUGU	2837
1374	CATCATTT C CATGGCTG	329	CAGCCAUG CUGAUGAG X CGAA AAAUGAUG	2838
1385	TGGCTGCT A GGCTGTGC	330	GCACAGCC CUGAUGAG X CGAA AGCAGCCA	2839
1406	AACTGGAT C CTACGCGG	331	CCGCGUAG CUGAUGAG X CGAA AUCCAGUU	2840
1409	TGGATCCT A CGCGGGAC	332	GUCCCGCG CUGAUGAG X CGAA AGGAUCCA	2841
1420	CGGGACGT C CTTTGTTT	333	AAACAAAG CUGAUGAG X CGAA ACGUCCCG	2842
1423	GACGTCCT T TGTTACG	334	CGUAAACA CUGAUGAG X CGAA AGGACGUC	2843
1424	ACGTCCTT T GTTACGT	335	ACGUAAAC CUGAUGAG X CGAA AAGGACGU	2844
1427	TCCTTTGT T TACGTCCC	336	GGGACGUA CUGAUGAG X CGAA ACAAAGGA	2845

Table 37

1428	CCTTTGTT T ACGTCCCG	337	CGGGACGU CUGAUGAG X CGAA AACAAAGG	2846
1429	CTTTGTTT A CGTCCCGT	338	ACGGGACG CUGAUGAG X CGAA AAACAAAG	2847
1433	GTTTACGT C CCGTCGGC	339	GCCGACGG CUGAUGAG X CGAA ACGUAAAC	2848
1438	CGTCCCGT C GCGCTGA	340	UCAGCGCC CUGAUGAG X CGAA ACGGGACG	2849
1449	CGCTGAAT C CCGCGGAC	341	GUCCGCGG CUGAUGAG X CGAA AUUCAGCG	2850
1465	CGACCCCT C CCGGGGCC	342	GGCCCCGG CUGAUGAG X CGAA AGGGGUCG	2851
1477	GGGCCGCT T GGGGCTCT	343	AGAGCCCC CUGAUGAG X CGAA AGCGGCC	2852
1484	TTGGGGCT C TACCGCCC	344	GGGCGGUA CUGAUGAG X CGAA AGCCCCAA	2853
1486	GGGGCTCT A CCGCCCGC	345	GCGGGCGG CUGAUGAG X CGAA AGAGCCCC	2854
1496	CGCCCGCT T CTCGCCT	346	AGGCGGAG CUGAUGAG X CGAA AGCGGGCG	2855
1497	GCCCGCTT C TCCGCTA	347	UAGGCGGA CUGAUGAG X CGAA AAGCGGGC	2856
1499	CCGCTTCT C CGCCTATT	348	AAUAGCGG CUGAUGAG X CGAA AGAAGCGG	2857
1505	CTCCGCTT A TTGTACCG	349	CGGUACAA CUGAUGAG X CGAA AGGCGGAG	2858
1507	CCGCTTAT T GTACCGAC	350	GUCGCUAC CUGAUGAG X CGAA AUAGGCGG	2859
1510	CCTATTGT A CCGACCGT	351	ACGUUCGG CUGAUGAG X CGAA ACAAUAGG	2860
1519	CCGACCGT C CACGGGGC	352	GCCCCGUG CUGAUGAG X CGAA ACGGUCGG	2861
1534	GCGCACCT C TCTTACG	353	CGUAAAGA CUGAUGAG X CGAA AGGUGCGC	2862
1536	GCACCTCT C TTTACGCG	354	CGCGUAAA CUGAUGAG X CGAA AGAGGUGC	2863
1538	ACCTCTCT T TACGCGGA	355	UCCGCGUA CUGAUGAG X CGAA AGAGAGGU	2864
1539	CCTCTCTT T ACGCGGAC	356	GUCCCGGU CUGAUGAG X CGAA AAGAGAGG	2865
1540	CTCTCTTT A CGCGGACT	357	AGUCCCGG CUGAUGAG X CGAA AAAGAGAG	2866
1549	CGCGGACT C CCCGCTG	358	CAGACGGG CUGAUGAG X CGAA AGUCCCGG	2867
1555	CTCCCCGT C TGTGCCTT	359	AAGGCACA CUGAUGAG X CGAA ACGGGGAG	2868
1563	CTGTGCCT T CTCATCTG	360	CAGAUGAG CUGAUGAG X CGAA AGGCACAG	2869
1564	TGTGCCTT C TCATCTGC	361	GCAGAUGA CUGAUGAG X CGAA AAGGCACA	2870
1566	TGCCTTCT C ATCTGCCG	362	CGGCAGAU CUGAUGAG X CGAA AGAAGGCA	2871
1569	CTTCTCAT C TGCCGGAC	363	GUCCGGCA CUGAUGAG X CGAA AUGAGAAG	2872
1588	TGTGCACT T CGCTTCAC	364	GUGAAGCG CUGAUGAG X CGAA AGUGCACA	2873
1589	GTGCACTT C GCTTCACC	365	GGUGAAGC CUGAUGAG X CGAA AAGUGCAC	2874
1593	ACTTCGCT T CACCTCTG	366	CAGAGGUG CUGAUGAG X CGAA AGCGAAGU	2875
1594	CTTCGCTT C ACCTCTGC	367	GCAGAGGU CUGAUGAG X CGAA AAGCGAAG	2876
1599	CTTCACCT C TGCACGTC	368	GACGUGCA CUGAUGAG X CGAA AGGUGAAG	2877
1607	CTGCACGT C GCATGGAG	369	CUCCAUGC CUGAUGAG X CGAA ACGUGCAG	2878
1651	CCCAAGGT C TTGCATAA	370	UUUAUGCA CUGAUGAG X CGAA ACCUUGGG	2879
1653	CAAGGTCT T GCATAAGA	371	UCUUAUGC CUGAUGAG X CGAA AGACCUUG	2880
1658	TCTTGCAT A AGAGGACT	372	AGUCCUCU CUGAUGAG X CGAA AUGCAAGA	2881
1667	AGAGGACT C TTGGACTT	373	AAGUCCAA CUGAUGAG X CGAA AGUCCUCU	2882
1669	AGGACTCT T GGACTTTC	374	GAAAGUCC CUGAUGAG X CGAA AGAGUCCU	2883
1675	CTTGGACT T TCAGCAAT	375	AUUGCUGA CUGAUGAG X CGAA AGUCCAAG	2884
1676	TTGGACTT T CAGCAATG	376	CAUUGCUG CUGAUGAG X CGAA AAGUCCAA	2885
1677	TGGACTTT C AGCAATGT	377	ACAUUGCUC CUGAUGAG X CGAA AAAGUCCA	2886
1686	AGCAATGT C AACGACCG	378	CGGUCGUU CUGAUGAG X CGAA ACAUUGCUC	2887
1699	ACCGACCT T GAGGCATA	379	UAUGCCUC CUGAUGAG X CGAA AGGUCGGU	2888
1707	TGAGGCAT A CTTCAAAG	380	CUUUGAAG CUGAUGAG X CGAA AUGCCUCA	2889
1710	GGCATACT T CAAAGACT	381	AGUCUUUG CUGAUGAG X CGAA AGUAUGCC	2890
1711	GCATACTT C AAAGACTG	382	CAGUCUUU CUGAUGAG X CGAA AAGUAUGC	2891
1725	CTGTGTGT T TAATGAGT	383	ACUCAUUA CUGAUGAG X CGAA ACACACAG	2892
1726	TGTGTGTT T AATGAGTG	384	CACUCAUU CUGAUGAG X CGAA AACACACA	2893
1727	GTGTGTTT A ATGAGTGG	385	CCACUCAU CUGAUGAG X CGAA AAACACAC	2894
1743	GGAGGAGT T GGGGGAGG	386	CCUCCCCC CUGAUGAG X CGAA ACUCCUCC	2895
1756	GAGGAGGT T AGGTTAAA	387	UUUAACCU CUGAUGAG X CGAA ACCUCCUC	2896

Table 37

1757	AGGAGGTT A GGTAAAG	388	CUUUAACC CUGAUGAG X CGAA AACCUCCU	2897
1761	GGTTAGGT T AAAGGTCT	389	AGACCUUU CUGAUGAG X CGAA ACCUAACC	2898
1762	GTTAGGTT A AAGGTCTT	390	AAGACCUU CUGAUGAG X CGAA AACCUAAC	2899
1768	TTAAAGGT C TTTGTACT	391	AGUACAAA CUGAUGAG X CGAA ACCUUUAA	2900
1770	AAAGGTCT T TGTACTAG	392	CUAGUACA CUGAUGAG X CGAA AGACCUUU	2901
1771	AAGGTCTT T GTACTAGG	393	CCUAGUAC CUGAUGAG X CGAA AAGACCUU	2902
1774	GTCTTTGT A CTAGGAGG	394	CCUCCUAG CUGAUGAG X CGAA ACAAAGAC	2903
1777	TTTGTACT A GGAGGCTG	395	CAGCCUCC CUGAUGAG X CGAA AGUACAAA	2904
1787	GAGGCTGT A GGCATAAA	396	UUUAUGCC CUGAUGAG X CGAA ACAGCCUC	2905
1793	GTAGGCAT A AATTGGTG	397	CACCAAUU CUGAUGAG X CGAA AUGCCUAC	2906
1797	GCATAAAT T GGTGTGTT	398	AACACACC CUGAUGAG X CGAA AUUUUAGC	2907
1805	TGGTGTGT T CACCAGCA	399	UGCUGGUG CUGAUGAG X CGAA ACACACCA	2908
1806	GGTGTGTT C ACCAGCAC	400	GUGCUGGU CUGAUGAG X CGAA AACACACC	2909
1824	ATGCAACT T TTTACCT	401	AGGUGAAA CUGAUGAG X CGAA AGUUGCAU	2910
1825	TGCAACTT T TTCACCTC	402	GAGGUGAA CUGAUGAG X CGAA AAGUUGCA	2911
1826	GCAACTTT T TCACCTCT	403	AGAGGUGA CUGAUGAG X CGAA AAAGUUGC	2912
1827	CAACTTTT T CACCTCTG	404	CAGAGGUG CUGAUGAG X CGAA AAAAGUUG	2913
1828	AACPTTTT C ACCTCTGC	405	GCAGAGGU CUGAUGAG X CGAA AAAAAGUU	2914
1833	TTTCACCT C TGCCTAAT	406	AUUAGGCA CUGAUGAG X CGAA AGGUGAAA	2915
1839	CTCTGCCT A ATCATCTC	407	GAGAUGAU CUGAUGAG X CGAA AGGCAGAG	2916
1842	TGCCTAAT C ATCTCATG	408	CAUGAGAU CUGAUGAG X CGAA AUUAGGCA	2917
1845	CTAATCAT C TCATGTTC	409	GAACAUGA CUGAUGAG X CGAA AUGAUUAG	2918
1847	AATCATCT C ATGTTCAT	410	AUGAACAU CUGAUGAG X CGAA AGAUGAUU	2919
1852	TCTCATGT T CATGTCCT	411	AGGACAUG CUGAUGAG X CGAA ACAUGAGA	2920
1853	CTCATGTT C ATGTCCTA	412	UAGGACAU CUGAUGAG X CGAA AACAUAGAG	2921
1858	GTTTCATGT C TACTGTGT	413	AACAGUAG CUGAUGAG X CGAA ACAUGAAC	2922
1861	CATGTCCT A CTGTTCAA	414	UUGACAG CUGAUGAG X CGAA AGGACAUG	2923
1866	CCTACTGT T CAAGCCTC	415	GAGGCUUG CUGAUGAG X CGAA ACAGUAGG	2924
1867	CTACTGTT C AAGCCTCC	416	GGAGGCUU CUGAUGAG X CGAA AACAGUAG	2925
1874	TCAAGCCT C CAAGCTGT	417	ACAGCUUG CUGAUGAG X CGAA AGGCUUGA	2926
1887	CTGTGCCT T GGGTGGCT	418	AGCCACCC CUGAUGAG X CGAA AGGCACAG	2927
1896	GGGTGGCT T TGGGGCAT	419	AUGCCCCA CUGAUGAG X CGAA AGCCACCC	2928
1897	GGTGGCTT T GGGGCATG	420	CAUGCCCC CUGAUGAG X CGAA AAGCCACC	2929
1911	ATGGACAT T GACCCGTA	421	UACGGGUC CUGAUGAG X CGAA AUGUCCAU	2930
1919	TGACCCGT A TAAAGAAT	422	AUUCUUUA CUGAUGAG X CGAA ACGGGUCA	2931
1921	ACCCGTAT A AAGAATTT	423	AAAUUCUU CUGAUGAG X CGAA AUACGGGU	2932
1928	TAAAGAAT T TGGAGCTT	424	AAGCUCCA CUGAUGAG X CGAA AUUCUUUA	2933
1929	AAAGAATT T GGAGCTTC	425	GAAGCUCC CUGAUGAG X CGAA AAUUCUUU	2934
1936	TTGGAGCT T CTGTGGAG	426	CUCCACAG CUGAUGAG X CGAA AGCUCCAA	2935
1937	TGGAGCTT C TGTGGAGT	427	ACUCCACA CUGAUGAG X CGAA AAGCUCCA	2936
1946	TGTGGAGT T ACTCTCTT	428	AAGAGAGU CUGAUGAG X CGAA ACUCCACA	2937
1947	GTGGAGTT A CTCTCTTT	429	AAAGAGAG CUGAUGAG X CGAA AACUCCAC	2938
1950	GAGTTACT C TCTTTTTT	430	AAAAAAGA CUGAUGAG X CGAA AGUAACUC	2939
1952	GTTACTCT C TTTTTCCT	431	GCAAAAAA CUGAUGAG X CGAA AGAGUAAAC	2940
1954	TACTCTCT T TTTTCCTT	432	AGGCAAAA CUGAUGAG X CGAA AGAGAGUA	2941
1955	ACTCTCTT T TTTGCCTT	433	AAGGCAAA CUGAUGAG X CGAA AAGAGAGU	2942
1956	CTCTCTTT T TTGCCTTC	434	GAAGGCAA CUGAUGAG X CGAA AAAGAGAG	2943
1957	TCTCTTTT T TGCTTCT	435	AGAAGGCA CUGAUGAG X CGAA AAAAGAGA	2944
1958	CTCTTTTT T GCCTTCTG	436	CAGAAGGC CUGAUGAG X CGAA AAAAAGAG	2945
1963	TTTTGCCT T CTGACTTC	437	GAAGUCAG CUGAUGAG X CGAA AGGCAAAA	2946
1964	TTTGCCTT C TGACTTCT	438	AGAAGUCA CUGAUGAG X CGAA AAGGCAAA	2947

Table 37

1970	TTCTGACT T CTTTCCTT	439	AAGGAAAG CUGAUGAG X CGAA AGUCAGAA	2948
1971	TCTGACTT C TTTCCTTC	440	GAAGGAAA CUGAUGAG X CGAA AAGUCAGA	2949
1973	TGACTTCT T TCCTTCTA	441	UAGAAGGA CUGAUGAG X CGAA AGAAGUCA	2950
1974	GACTTCTT T CCTTCTAT	442	AUAGAAGG CUGAUGAG X CGAA AAGAAGUC	2951
1975	ACTTCTTT C CTTCTATT	443	AAUAGAAG CUGAUGAG X CGAA AAAGAAGU	2952
1978	TCTTTCCT T CTATTCGA	444	UCGAAUAG CUGAUGAG X CGAA AGGAAAGA	2953
1979	CTTTCCTT C TATTCGAG	445	CUCGAAUA CUGAUGAG X CGAA AAGGAAAG	2954
1981	TTCTTCTT A TTCGAGAT	446	AUCUCGAA CUGAUGAG X CGAA AGAAGGAA	2955
1983	CCTTCTAT T CGAGATCT	447	AGAUCUCG CUGAUGAG X CGAA AUAGAAGG	2956
1984	CTTCTATT C GAGATCTC	448	GAGAUCUC CUGAUGAG X CGAA AAUAGAAG	2957
1990	TTCGAGAT C TCCTCGAC	449	GUCGAGGA CUGAUGAG X CGAA AUCUCGAA	2958
1992	CGAGATCT C CTCGACAC	450	GUGUCGAG CUGAUGAG X CGAA AGAUCUCG	2959
1995	GATCTCCT C GACACCGC	451	GCGGUGUC CUGAUGAG X CGAA AGGAGAUC	2960
2006	CACCGCCT C TGCTCTGT	452	ACAGAGCA CUGAUGAG X CGAA AGGCGGUG	2961
2011	CCTCTGCT C TGTATCGG	453	CCGAUACA CUGAUGAG X CGAA AGCAGAGG	2962
2015	TGCTCTGT A TCGGGGGG	454	CCCCCGCA CUGAUGAG X CGAA ACAGAGCA	2963
2017	CTCTGTAT C GGGGGGCC	455	GGCCCCC CUGAUGAG X CGAA AUACAGAG	2964
2027	GGGGGCCT T AGAGTCTC	456	GAGACUCU CUGAUGAG X CGAA AGGCCCCC	2965
2028	GGGGCCTT A GAGTCTCC	457	GGAGACUC CUGAUGAG X CGAA AAGGCCCC	2966
2033	CTTAGAGT C TCCGGAAC	458	GUUCCGGA CUGAUGAG X CGAA ACUCUAAG	2967
2035	TAGAGTCT C CGGAACAT	459	AUGUCCG CUGAUGAG X CGAA AGACUCUA	2968
2044	CGGAACAT T GTTCACCT	460	AGGUGAAC CUGAUGAG X CGAA AUGUCCG	2969
2047	AACATTGT T CACCTCAC	461	GUGAGGUG CUGAUGAG X CGAA ACAAUGUU	2970
2048	ACATTGTT C ACCTCACC	462	GGUGAGGU CUGAUGAG X CGAA AACAAUGU	2971
2053	GTTACCTT C ACCATACG	463	CGUAUGGU CUGAUGAG X CGAA AGGUGAAC	2972
2059	CTCACCAT A CGGCACTC	464	GAGUGCCG CUGAUGAG X CGAA AUGGUGAG	2973
2067	ACGGCACT C AGGCAAGC	465	GCUUGCCU CUGAUGAG X CGAA AGUGCCGU	2974
2077	GGCAAGCT A TTCTGTGT	466	ACACAGAA CUGAUGAG X CGAA AGCUUGCC	2975
2079	CAAGCTAT T CTGTGTTG	467	CAACACAG CUGAUGAG X CGAA AUAGCUUG	2976
2080	AAGCTATT C TGTGTTGG	468	CCAACACA CUGAUGAG X CGAA AAUAGCUU	2977
2086	TTCTGTGT T GGGGTGAG	469	CUCACCCC CUGAUGAG X CGAA ACACAGAA	2978
2096	GGGTGAGT T GATGAATC	470	GAUUCAUC CUGAUGAG X CGAA ACUCACCC	2979
2104	TGATGAAT C TAGCCACC	471	GGUGGCUA CUGAUGAG X CGAA AUUCAUCA	2980
2106	ATGAATCT A GCCACCTG	472	CAGGUGGC CUGAUGAG X CGAA AGAUUCAU	2981
2125	TGGGAAGT A ATTTGGAA	473	UUCCAAU CUGAUGAG X CGAA ACUUCCCA	2982
2128	GAAGTAAT T TGGAAGAT	474	AUCUCCA CUGAUGAG X CGAA AUUACUUC	2983
2129	AAGTAATT T GGAAGATC	475	GAUCUUC CUGAUGAG X CGAA AAUUAUUC	2984
2137	TGGAAGAT C CAGCATCC	476	GGAUGCUG CUGAUGAG X CGAA AUCUCCA	2985
2144	TCCAGCAT C CAGGGAAT	477	AUUCUUG CUGAUGAG X CGAA AUGCUGGA	2986
2153	CAGGGAAT T AGTAGTCA	478	UGACUACU CUGAUGAG X CGAA AUUCCUG	2987
2154	AGGGAATT A GTAGTCAG	479	CUGACUAC CUGAUGAG X CGAA AAUUCUUC	2988
2157	GAATTAGT A GTCAGCTA	480	UAGCUGAC CUGAUGAG X CGAA ACUAAUUC	2989
2160	TTAGTAGT C AGCTATGT	481	ACAUAGCU CUGAUGAG X CGAA ACUACUAA	2990
2165	AGTCAGCT A TGCAACG	482	CGUUGACA CUGAUGAG X CGAA AGCUGACU	2991
2169	AGCTATGT C AACGTTAA	483	UUAACGUU CUGAUGAG X CGAA ACAUAGCU	2992
2175	GTCAACGT T AATATGGG	484	CCCAUAUU CUGAUGAG X CGAA ACGUUGAC	2993
2176	TCAACGTT A ATATGGGC	485	GCCCAUAU CUGAUGAG X CGAA AACGUUGA	2994
2179	ACGTTAAT A TGGGCCTA	486	UAGGCCCA CUGAUGAG X CGAA AUUAACGU	2995
2187	ATGGGCCT A AAAATCAG	487	CUGAUUUU CUGAUGAG X CGAA AGGCCCAU	2996
2193	CTAAAAAT C AGACAAC	488	AGUUGUCU CUGAUGAG X CGAA AUUUUAG	2997
2202	AGACAAC T TGTGTTT	489	AACCACAA CUGAUGAG X CGAA AGUUGUCU	2998

Table 57

2204	ACAACTAT T GTGGTTTC	490	GAAACCAC CUGAUGAG X CGAA AUAGUUGU	2999
2210	ATTGTGGT T TCACATTT	491	AAAUGUGA CUGAUGAG X CGAA ACCACAAU	3000
2211	TTGTGGTT T CACATTTTC	492	GAAAUGUG CUGAUGAG X CGAA AACCACAA	3001
2212	TGTGGTTT C ACATTTCC	493	GGAAAUGU CUGAUGAG X CGAA AAACCACA	3002
2217	TTTCACAT T TCCTGTCT	494	AGACAGGA CUGAUGAG X CGAA AUGUGAAA	3003
2218	TTACATT T CCTGTCTT	495	AAGACAGG CUGAUGAG X CGAA AAUGUGAA	3004
2219	TCACATTT C CTGTCTTA	496	UAAGACAG CUGAUGAG X CGAA AAAUGUGA	3005
2224	TTTCCTGT C TTACTTTT	497	AAAAGUAA CUGAUGAG X CGAA ACAGGAAA	3006
2226	TCCTGTCT T ACTTTTGG	498	CCAAAAGU CUGAUGAG X CGAA AGACAGGA	3007
2227	CCTGTCTT A CTTTGGG	499	CCCAAAAG CUGAUGAG X CGAA AAGACAGG	3008
2230	GTCTTACT T TTGGGCGA	500	UCGCCCAA CUGAUGAG X CGAA AGUAAGAC	3009
2231	TCTTACTT T TGGGCGAG	501	CUCGCCCA CUGAUGAG X CGAA AAGUAAGA	3010
2232	CTTACTTT T GGGCGAGA	502	UCUCGCCC CUGAUGAG X CGAA AAAGUAAG	3011
2247	GAAACTGT T CTTGAATA	503	UAUUCAG CUGAUGAG X CGAA ACAGUUUC	3012
2248	AAACTGTT C TTGAATAT	504	AUAUCAA CUGAUGAG X CGAA AACAGUUU	3013
2250	ACTGTTCT T GAATATTT	505	AAAUUUC CUGAUGAG X CGAA AGAACAGU	3014
2255	TCTTGAAT A TTTGGTGT	506	ACACCAAA CUGAUGAG X CGAA AUUCAAGA	3015
2257	TTGAATAT T TGGTGTCT	507	AGACACCA CUGAUGAG X CGAA AUAUUCAA	3016
2258	TGAATATT T GGTGTCTT	508	AAGACACC CUGAUGAG X CGAA AAUAUUCA	3017
2264	TTTGGTGT C TTTTGGAG	509	CUCCAAAA CUGAUGAG X CGAA ACACCAAA	3018
2266	TGGTGTCT T TTGGAGTG	510	CACUCCAA CUGAUGAG X CGAA AGACACCA	3019
2267	GGTGTCTT T TGGAGTGT	511	ACACUCCA CUGAUGAG X CGAA AAGACACC	3020
2268	GTGTCTTT T GGAGTGTG	512	CACACUCC CUGAUGAG X CGAA AAAGACAC	3021
2280	GTGTGGAT T CGCACTCC	513	GGAGUGCG CUGAUGAG X CGAA AUCCACAC	3022
2281	TGTGGATT C GCACTCCT	514	AGGAGUGC CUGAUGAG X CGAA AAUCCACA	3023
2287	TTGCACT C CTCCTGCA	515	UGCAGGAG CUGAUGAG X CGAA AGUGCGAA	3024
2290	GCACTCCT C CTGCATAT	516	AUAUGCAG CUGAUGAG X CGAA AGGAGUGC	3025
2297	TCCTGCAT A TAGACCAC	517	GUGGUCUA CUGAUGAG X CGAA AUGCAGGA	3026
2299	CTGCATAT A GACCACCA	518	UGGUGGUC CUGAUGAG X CGAA AUAUGCAG	3027
2317	ATGCCCT A TCTTATCA	519	UGAUAAGA CUGAUGAG X CGAA AGGGGCAU	3028
2319	GCCCTAT C TTATCAAC	520	GUUGAUAA CUGAUGAG X CGAA AUAGGGGC	3029
2321	CCCTATCT T ATCAACAC	521	GUGUUGAU CUGAUGAG X CGAA AGAUAGGG	3030
2322	CCTATCTT A TCAACACT	522	AGUGUUGA CUGAUGAG X CGAA AAGAUAGG	3031
2324	TATCTTAT C AACACTTC	523	GAAGUGUU CUGAUGAG X CGAA AUAAGAU	3032
2331	TCAACACT T CCGGAAAC	524	GUUUCGG CUGAUGAG X CGAA AGUGUUGA	3033
2332	CAACACTT C CGGAAACT	525	AGUUUCCG CUGAUGAG X CGAA AAGUGUUG	3034
2341	CGGAAACT A CTGTGTGT	526	AACAACAG CUGAUGAG X CGAA AGUUUCCG	3035
2346	ACTACTGT T GTTAGACG	527	CGUCUAAC CUGAUGAG X CGAA ACAGUAGU	3036
2349	ACTGTTGT T AGACGAAG	528	CUUCGUCU CUGAUGAG X CGAA ACAACAGU	3037
2350	CTGTTGTT A GACGAAGA	529	UCUUCGUC CUGAUGAG X CGAA AACACAG	3038
2366	AGGCAGGT C CCCTAGAA	530	UUCUAGGG CUGAUGAG X CGAA ACCUGCCU	3039
2371	GGTCCCCT A GAAGAAGA	531	UCUUCUUC CUGAUGAG X CGAA AGGGGACC	3040
2383	GAAGAACT C CCTCGCCT	532	AGGCGAGG CUGAUGAG X CGAA AGUUCUUC	3041
2387	AACTCCCT C GCCTCGCA	533	UGCAGGCG CUGAUGAG X CGAA AGGGAGUU	3042
2392	CCTCGCCT C GCAGACGA	534	UCGUCUGC CUGAUGAG X CGAA AGGCGAGG	3043
2405	ACGAAGGT C TCAATCGC	535	GCGAUUGA CUGAUGAG X CGAA ACCUUCGU	3044
2407	GAAGGTCT C AATCGCCG	536	CGGCGAUU CUGAUGAG X CGAA AGACCUUC	3045
2411	GTCTCAAT C GCCGCGTC	537	GACGCGGC CUGAUGAG X CGAA AUUGAGAC	3046
2419	CGCCGCGT C GCAGAAGA	538	UCUUCGCG CUGAUGAG X CGAA ACGCGGCG	3047
2429	CAGAAGAT C TCAATCTC	539	GAGAUUGA CUGAUGAG X CGAA AUCUUCUG	3048
2431	GAAGATCT C AATCTCGG	540	CCGAGAUU CUGAUGAG X CGAA AGAUCUUC	3049

Table 37

2435	ATCTCAAT C TCGGAAT	541	AUUCCCGA CUGAUGAG X CGAA AUUGAGAU	3050
2437	CTCAATCT C GGAATCT	542	AGAUUCCC CUGAUGAG X CGAA AGAUUGAG	3051
2444	TCGGGAAT C TCAATGTT	543	AACAUGA CUGAUGAG X CGAA AUUCCCGA	3052
2446	GGAATCT C AATGTTAG	544	CUAACAUU CUGAUGAG X CGAA AGAUUCCC	3053
2452	CTCAATGT T AGTATTCC	545	GGAAUACU CUGAUGAG X CGAA ACAUUGAG	3054
2453	TCAATGTT A GTATTCCT	546	AGGAAUAC CUGAUGAG X CGAA AACAUUGA	3055
2456	ATGTTAGT A TTCCTTGG	547	CCAAGGAA CUGAUGAG X CGAA ACUAACAU	3056
2458	GTTAGTAT T CCTTGGAC	548	GUCCAAGG CUGAUGAG X CGAA AUACU AAC	3057
2459	TTAGTATT C CTTGACAC	549	UGUCCAAG CUGAUGAG X CGAA AAUACUAA	3058
2462	GTATTCCT T GGACACAT	550	AUGUGUCC CUGAUGAG X CGAA AGGAAUAC	3059
2471	GGACACAT A AGGTGGGA	551	UCCCACCU CUGAUGAG X CGAA AUGUGUCC	3060
2484	GGGAAACT T TACGGGGC	552	GCCCCGUA CUGAUGAG X CGAA AGUUUCCC	3061
2485	GGAAACTT T ACGGGGCT	553	AGCCCCGU CUGAUGAG X CGAA AAGUUUCC	3062
2486	GAAACTTT A CGGGGCTT	554	AAGCCCCG CUGAUGAG X CGAA AAAGUUUC	3063
2494	ACGGGGCT T TATTCTTC	555	GAAGAAUA CUGAUGAG X CGAA AGCCCCGU	3064
2495	CGGGGCTT T ATTCTTCT	556	AGAAGAAU CUGAUGAG X CGAA AAGCCCCG	3065
2496	GGGGCTTT A TTCTTCTA	557	UAGAAGAA CUGAUGAG X CGAA AAAGCCCC	3066
2498	GGCTTTAT T CTTCTACG	558	CGUAGAAG CUGAUGAG X CGAA AUAAAGCC	3067
2499	GCTTTATT C TTCTACGG	559	CCGUAGAA CUGAUGAG X CGAA AAUAAAGC	3068
2501	TTTATTCT T CTACGGTA	560	UACCGUAG CUGAUGAG X CGAA AGAAUAAA	3069
2502	TTATTCTT C TACGGTAC	561	GUACCGUA CUGAUGAG X CGAA AAGAAUAA	3070
2504	ATTCTTCT A CGGTACCT	562	AGGUACCG CUGAUGAG X CGAA AGAAGAAU	3071
2509	TCTACGGT A CCTTGCTT	563	AAGCAAGG CUGAUGAG X CGAA ACCGUAGA	3072
2513	CGGTACCT T GCTTTAAT	564	AUUAAAGC CUGAUGAG X CGAA AGGUACCG	3073
2517	ACCTTGCT T TAATCCTA	565	UAGGAUUA CUGAUGAG X CGAA AGCAAGGU	3074
2518	CCTTGCTT T AATCCTAA	566	UUAGGAUU CUGAUGAG X CGAA AAGCAAGG	3075
2519	CTTGCTTT A ATCCTAAA	567	UUUAGGAU CUGAUGAG X CGAA AAAGCAAG	3076
2522	GCTTTAAT C CTAAATGG	568	CCAUUUAG CUGAUGAG X CGAA AUUAAAGC	3077
2525	TTAATCCT A AATGGCAA	569	UUGCCAUU CUGAUGAG X CGAA AGGAUUAA	3078
2537	GGCAAAC C CTTCTTTT	570	AAAAGAAG CUGAUGAG X CGAA AGUUUGCC	3079
2540	AAACTCCT T CTTTTCCT	571	AGGAAAAG CUGAUGAG X CGAA AGGAGUUU	3080
2541	AACTCCTT C TTTTCTTG	572	CAGGAAAA CUGAUGAG X CGAA AAGGAGUU	3081
2543	CTCCTTCT T TTCCTGAC	573	GUCAGGAA CUGAUGAG X CGAA AGAAGGAG	3082
2544	TCCTTCTT T TCCTGACA	574	UGUCAGGA CUGAUGAG X CGAA AAGAAGGA	3083
2545	CCTTCTTT T CCTGACAT	575	AUGUCAGG CUGAUGAG X CGAA AAAGAAGG	3084
2546	CTTCTTTT C CTGACATT	576	AAUGUCAG CUGAUGAG X CGAA AAAAGAAG	3085
2554	CCTGACAT T CATTTGCA	577	UGCAAAUG CUGAUGAG X CGAA AUGUCAGG	3086
2555	CTGACATT C ATTTGCAG	578	CUGCAAAU CUGAUGAG X CGAA AAUGUCAG	3087
2558	ACATTCAT T TGCAGGAG	579	CUCCUGCA CUGAUGAG X CGAA AUGAAUGU	3088
2559	CATTCATT T GCAGGAGG	580	CCUCCUGC CUGAUGAG X CGAA AAUGAAUG	3089
2572	GAGGACAT T GTTGATAG	581	CUAUCAAC CUGAUGAG X CGAA AUGUCCUC	3090
2575	GACATTGT T GATAGATG	582	CAUCUAUC CUGAUGAG X CGAA ACAAUGUC	3091
2579	TTGTTGAT A GATGTAAG	583	CUUACAUC CUGAUGAG X CGAA AUCAACAA	3092
2585	ATAGATGT A AGCAATTT	584	AAAUUGCU CUGAUGAG X CGAA ACAUCUUA	3093
2592	TAAGCAAT T TGTGGGGC	585	GCCCCACA CUGAUGAG X CGAA AUUGCUUA	3094
2593	AAGCAATT T GTGGGGCC	586	GGCCCCAC CUGAUGAG X CGAA AAUUGCUU	3095
2605	GGGCCCTT T ACAGTAAA	587	UUUACUGU CUGAUGAG X CGAA AGGGGGCC	3096
2606	GGCCCCTT A CAGTAAAT	588	AUUUACUG CUGAUGAG X CGAA AAGGGGGC	3097
2611	CTTACAGT A AATGAAAA	589	UUUUAUUU CUGAUGAG X CGAA ACUGUAAG	3098
2629	AGGAGACT T AAATTAAC	590	GUUAAUUU CUGAUGAG X CGAA AGUCUCCU	3099
2630	GGAGACTT A AATTAAC	591	AGUUAUUU CUGAUGAG X CGAA AAGUCUCC	3100

Table 37

2634	ACTTAAAT T AACTATGC	592	GCAUAGUU CUGAUGAG X CGAA AUUUAAGU	3101
2635	CTTAAATT A ACTATGCC	593	GGCAUAGU CUGAUGAG X CGAA AAUUAAG	3102
2639	AATTAACT A TGCCTGCT	594	AGCAGGCA CUGAUGAG X CGAA AGUUAUU	3103
2648	TGCCTGCT A GGTTTTAT	595	AUAAAACC CUGAUGAG X CGAA AGCAGGCA	3104
2652	TGCTAGGT T TTATCCCA	596	UGGGAUAA CUGAUGAG X CGAA ACCUAGCA	3105
2653	GCTAGGTT T TATCCCAA	597	UUGGGAUA CUGAUGAG X CGAA AACCUAGC	3106
2654	CTAGGTTT T ATCCCAAT	598	AUUGGGAU CUGAUGAG X CGAA AAACCUAG	3107
2655	TAGGTTTT A TCCCAATG	599	CAUUGGGA CUGAUGAG X CGAA AAAACCUA	3108
2657	GGTTTTAT C CCAATGTT	600	AACAUUGG CUGAUGAG X CGAA AUAAAACC	3109
2665	CCCAATGT T ACTAAATA	601	UAUUUAGU CUGAUGAG X CGAA ACAUUGGG	3110
2666	CCAATGTT A CTAAATAT	602	AUAUUUAG CUGAUGAG X CGAA AACAUUGG	3111
2669	ATGTTACT A AATATTTG	603	CAAAUAUU CUGAUGAG X CGAA AGUAACAU	3112
2673	TACTAAAT A TTTGCCCT	604	AGGGCAAA CUGAUGAG X CGAA AUUUAGUA	3113
2675	CTAAATAT T TGCCTTA	605	UAAGGGCA CUGAUGAG X CGAA AUUUUAG	3114
2676	TAAATATT T GCCCTTAG	606	CUAAGGGC CUGAUGAG X CGAA AAUAUUUA	3115
2682	TTTGCCCT T AGATAAAG	607	CUUUUUCU CUGAUGAG X CGAA AGGGCAAA	3116
2683	TTGCCCTT A GATAAAGG	608	CCUUUAUC CUGAUGAG X CGAA AAGGGCAA	3117
2687	CCTTAGAT A AAGGGATC	609	GAUCCCUU CUGAUGAG X CGAA AUCUAAGG	3118
2695	AAAGGGAT C AAACCGTA	610	UACGGUUU CUGAUGAG X CGAA AUCCCUUU	3119
2703	CAAACCGT A TTATCCAG	611	CUGGAUAA CUGAUGAG X CGAA ACGGUUUG	3120
2705	AACCGTAT T ATCCAGAG	612	CUCUGGAU CUGAUGAG X CGAA AUACGGUU	3121
2706	ACCGTATT A TCCAGAGT	613	ACUCUGGA CUGAUGAG X CGAA AAUACGGU	3122
2708	CGTATTAT C CAGAGTAT	614	AUACUCUG CUGAUGAG X CGAA AUAAUACG	3123
2715	TCCAGAGT A TGTAGTTA	615	UAACUACA CUGAUGAG X CGAA ACUCUGGA	3124
2719	GAGTATGT A GTTAATCA	616	UGAUUAAC CUGAUGAG X CGAA ACAUACUC	3125
2722	TATGTAGT T AATCATT A	617	UAAUGAUU CUGAUGAG X CGAA ACUACUA	3126
2723	ATGTAGTT A ATCATTAC	618	GUAAUGAU CUGAUGAG X CGAA AACUACAU	3127
2726	TAGTTAAT C ATTACTTC	619	GAAGUAAU CUGAUGAG X CGAA AUUAACUA	3128
2729	TTAATCAT T ACTTCCAG	620	CUGGAAGU CUGAUGAG X CGAA AUGAUUAA	3129
2730	TAATCATT A CTTCAG A	621	UCUGGAAG CUGAUGAG X CGAA AAUGAUUA	3130
2733	TCATTACT T CCAGACGC	622	GCGUCUGG CUGAUGAG X CGAA AGUAAUGA	3131
2734	CATTACTT C CAGACGCG	623	CGCGUCUG CUGAUGAG X CGAA AAGUAAUG	3132
2747	CGCGACAT T ATTTACAC	624	GUGUAAAU CUGAUGAG X CGAA AUGUCGCG	3133
2748	GCGACATT A TTTACACA	625	UGUGUAAA CUGAUGAG X CGAA AAUGUCGC	3134
2750	GACATTAT T TACACACT	626	AGUGUGUA CUGAUGAG X CGAA AUAAUGUC	3135
2751	ACATTATT T ACACACTC	627	GAGUGUGU CUGAUGAG X CGAA AAUAAUGU	3136
2752	CATTATTT A CACACTCT	628	AGAGUGUG CUGAUGAG X CGAA AAAUAAUG	3137
2759	TACACACT C TTTGGAAG	629	CUUCCAAA CUGAUGAG X CGAA AGUGUGUA	3138
2761	CACACTCT T TGGAAGGC	630	GCCUCCAA CUGAUGAG X CGAA AGAGUGUG	3139
2762	ACACTCTT T GGAAGGCG	631	CGCCUCC CUGAUGAG X CGAA AAGAGUGU	3140
2776	GCGGGGAT C TTATATAA	632	UUUAUUA CUGAUGAG X CGAA AUCCCGC	3141
2778	GGGATCT T ATATAAAA	633	UUUUUAU CUGAUGAG X CGAA AGAUCCC	3142
2779	GGGATCTT A TATAAAG	634	CUUUUAU CUGAUGAG X CGAA AAGAUC	3143
2781	GATCTTAT A TAAAGAG	635	CUCUUUUA CUGAUGAG X CGAA AUAAGAUC	3144
2783	TCTTATAT A AAAGAGAG	636	CUCUCUUU CUGAUGAG X CGAA AUAUAAGA	3145
2793	AAGAGAGT C CACACGTA	637	UACGUGUG CUGAUGAG X CGAA ACUCUCUU	3146
2801	CCACACGT A GCGCCTCA	638	UGAGGCGC CUGAUGAG X CGAA ACGUGUGG	3147
2808	TAGCGCCT C ATTTTGCG	639	CGCAAAAU CUGAUGAG X CGAA AGGCGCUA	3148
2811	CGCCTCAT T TTGCGGGT	640	ACCCGCAA CUGAUGAG X CGAA AUGAGGCG	3149
2812	GCCTCATT T TGCGGGTC	641	GACCCGCA CUGAUGAG X CGAA AAUGAGGC	3150
2813	CCTCATTT T GCGGGTCA	642	UGACCCGC CUGAUGAG X CGAA AAAUGAGG	3151

Table 37

2820	TTGCGGGT C ACCATATT	643	AAUAUGGU CUGAUGAG X CGAA ACCCGCAA	3152
2826	GTCACCAT A TTCTTGGG	644	CCCAAGAA CUGAUGAG X CGAA AUGGUGAC	3153
2828	CACCATAT T CTTGGGAA	645	UUCCCAAG CUGAUGAG X CGAA AUAUGGUG	3154
2829	ACCATATT C TTGGGAAC	646	GUUCCCAA CUGAUGAG X CGAA AAUAUGGU	3155
2831	CATATTCT T GGGAACAA	647	UUGUCCCC CUGAUGAG X CGAA AGAAUAUG	3156
2843	AACAAGAT C TACAGCAT	648	AUGCUGUA CUGAUGAG X CGAA AUCUUGUU	3157
2845	CAAGATCT A CAGCATGG	649	CCAUGCUG CUGAUGAG X CGAA AGAUCUUG	3158
2859	TGGGAGGT T GGTCTTCC	650	GGAAAGCC CUGAUGAG X CGAA ACCUCCCA	3159
2863	AGGTTGGT C TTCAAAC	651	GUUUGGAA CUGAUGAG X CGAA ACCAACCU	3160
2865	GTTGGTCT T CCAAACCT	652	AGGUUUGG CUGAUGAG X CGAA AGACCAAC	3161
2866	TTGGTCTT C CAAACCTC	653	GAGGUUUG CUGAUGAG X CGAA AAGACCAA	3162
2874	CCAAACCT C GAAAAGGC	654	GCCUUUUC CUGAUGAG X CGAA AGGUUUGG	3163
2895	GGACAAAT C TTTGTGTC	655	GACAGAAA CUGAUGAG X CGAA AUUUGUCC	3164
2897	ACAAATCT T TCTGTCCC	656	GGGACAGA CUGAUGAG X CGAA AGAUUUGU	3165
2898	CAAATCTT T CTGTCCCC	657	GGGGACAG CUGAUGAG X CGAA AAGAUUUG	3166
2899	AAATCTTT C TGTCCTCA	658	UGGGGACA CUGAUGAG X CGAA AAAGAUUU	3167
2903	CTTCTGT C CCAATCC	659	GGAUUGGG CUGAUGAG X CGAA ACAGAAAG	3168
2910	TCCCCAAT C CCCTGGGA	660	UCCCAGGG CUGAUGAG X CGAA AUUGGGGA	3169
2920	CCTGGGAT T CTTCCTCG	661	CGGGGAAG CUGAUGAG X CGAA AUCCCAGG	3170
2921	CTGGGATT C TTCCCCGA	662	UCGGGGAA CUGAUGAG X CGAA AAUCCAG	3171
2923	GGGATTCT T CCCCAGTC	663	GAUCGGG CUGAUGAG X CGAA AGAAUCCC	3172
2924	GGATTCTT C CCCGATCA	664	UGAUCGGG CUGAUGAG X CGAA AAGAAUCC	3173
2931	TCCCCGAT C ATCAGTTG	665	CAACUGAU CUGAUGAG X CGAA AUCGGGGA	3174
2934	CCGATCAT C AGTTGGAC	666	GUCCAACU CUGAUGAG X CGAA AUGAUCGG	3175
2938	TCATCAGT T GGACCTG	667	CAGGGUCC CUGAUGAG X CGAA ACUGAUGA	3176
2950	CCCTGCAT T CAAAGCCA	668	UGGCUUUG CUGAUGAG X CGAA AUGCAGGG	3177
2951	CCTGCATT C AAAGCCAA	669	UUGGCUUU CUGAUGAG X CGAA AAUGCAGG	3178
2962	AGCCAAT C AGTAAATC	670	GAUUUACU CUGAUGAG X CGAA AGUUGGCU	3179
2966	AACTCAGT A AATCCAGA	671	UCUGGAU CUGAUGAG X CGAA ACUGAGUU	3180
2970	CAGTAAAT C CAGATTGG	672	CCAUCUG CUGAUGAG X CGAA AUUUACUG	3181
2976	ATCCAGAT T GGGACCTC	673	GAGGUCCC CUGAUGAG X CGAA AUCUGGAU	3182
2984	TGGGACCT C AACC CGCA	674	UGCGGGUU CUGAUGAG X CGAA AGGUCCCA	3183
3037	GGGAGCAT T CGGGCCAG	675	CUGGCCCC CUGAUGAG X CGAA AUGCUCCC	3184
3038	GGAGCATT C GGGCCAGG	676	CCUGGCCC CUGAUGAG X CGAA AAUGCUC	3185
3049	GCCAGGGT T CACCCCTC	677	GAGGGGUG CUGAUGAG X CGAA ACCCUGGC	3186
3050	CCAGGGT C ACCCCTCC	678	GGAGGGGU CUGAUGAG X CGAA AACCCUGG	3187
3057	TCACCCCT C CCCATGGG	679	CCCAUGGG CUGAUGAG X CGAA AGGGGUGA	3188
3073	GGGACTGT T GGGGTGGA	680	UCCACCCC CUGAUGAG X CGAA ACAGUCCC	3189
3087	GGAGCCCT C ACGCTCAG	681	CUGAGCGU CUGAUGAG X CGAA AGGGCUCC	3190
3093	CTCACGCT C AGGGCCTA	682	UAGGCCCU CUGAUGAG X CGAA AGCUGAG	3191
3101	CAGGGCCT A CTCACAAC	683	GUUGUGAG CUGAUGAG X CGAA AGGCCUG	3192
3104	GGCCTACT C ACAACTGT	684	ACAGUUGU CUGAUGAG X CGAA AGUAGGCC	3193
3123	CAGCAGCT C CTCCTCCT	685	AGGAGGAG CUGAUGAG X CGAA AGCUGCUG	3194
3126	CAGCTCCT C CTCCTGCC	686	GGCAGGAG CUGAUGAG X CGAA AGGAGCUG	3195
3129	CTCCTCCT C CTGCCTCC	687	GGAGGCAG CUGAUGAG X CGAA AGGAGGAG	3196
3136	TCCTGCCT C CACCAATC	688	GAUUGGUG CUGAUGAG X CGAA AGGCAGGA	3197
3144	CCACCAAT C GGCAGTCA	689	UGACUGCC CUGAUGAG X CGAA AUUGGUGG	3198
3151	TCGGCAGT C AGGAAGGC	690	GCCUCCU CUGAUGAG X CGAA ACUGCCGA	3199
3165	GGCAGCCT A CTCCTTA	691	UAAGGGAG CUGAUGAG X CGAA AGGCUGCC	3200
3168	AGCCTACT C CCTTATCT	692	AGAUAAAG CUGAUGAG X CGAA AGUAGGCU	3201
3172	TACTCCCT T ATCTCCAC	693	GUGGAGAU CUGAUGAG X CGAA AGGGAGUA	3202

Table 3/

3173	ACTCCCTT A TCTCCACC	694	GGUGGAGA CUGAUGAG X CGAA AAGGGAGU	3203
3175	TCCCTTAT C TCCACCTC	695	GAGGUGGA CUGAUGAG X CGAA AUAAGGGA	3204
3177	CCTTATCT C CACCTCTA	696	UAGAGGUG CUGAUGAG X CGAA AGAUAAGG	3205
3183	CTCCACCT C TAAGGGAC	697	GUCCCUUA CUGAUGAG X CGAA AGGUGGAG	3206
3185	CCACCTCT A AGGGACAC	698	GUGUCCCU CUGAUGAG X CGAA AGAGGUGG	3207
3195	GGGACACT C ATCCTCAG	699	CUGAGGAU CUGAUGAG X CGAA AGUGUCCC	3208
3198	ACACTCAT C CTCAGGCC	700	GGCCUGAG CUGAUGAG X CGAA AUGAGUGU	3209
3201	CTCATCCT C AGGCCATG	701	CAUGGCCU CUGAUGAG X CGAA AGGAUGAG	3210

Input Sequence = AF100308. Cut Site = UH/.

Stem Length = 8 . Core Sequence = CUGAUGAG X CGAA (X = GCCGUUAGGC or other stem II)

AF100308 (Hepatitis B virus strain 2-18, 3215 bp)

Table 38

Table 38: Human HBV Inozyme and Substrate Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
9	AACTCCAC C ACTTTCCA	702	UGGAAAGU CUGAUGAG X CGAA IUGGAGUU	3211
10	ACTCCACC A CTTTCAC	703	GUGGAAAG CUGAUGAG X CGAA IGUGGAGU	3212
12	TCCACCAC T TTCCACCA	704	UGGUGGAA CUGAUGAG X CGAA IUGGUGGA	3213
16	CCACTTTC C ACCAAACT	705	AGUUUGGU CUGAUGAG X CGAA IAAAGUGG	3214
17	CACTTTCC A CCAAATC	706	GAGUUUGG CUGAUGAG X CGAA IGAAAGUG	3215
19	CTTTCCAC C AAATCTT	707	AAGAGUUU CUGAUGAG X CGAA IUGGAAAG	3216
20	TTTCCACC A AATCTTC	708	GAAGAGUU CUGAUGAG X CGAA IGUGGAAA	3217
24	CACCAAAC T CTTCAAGA	709	UCUUGAAG CUGAUGAG X CGAA IUUUGGUG	3218
26	CCAAACTC T TCAAGATC	710	GAUCUUGA CUGAUGAG X CGAA IAGUUUGG	3219
29	AACTCTTC A AGATCCCA	711	UGGGAUCU CUGAUGAG X CGAA IAAGAGUU	3220
35	TCAAGATC C CAGATCA	712	UGACUCUG CUGAUGAG X CGAA IAUUCUGA	3221
36	CAAGATCC C AGATCAG	713	CUGACUCU CUGAUGAG X CGAA IGAUCUUG	3222
37	AAGATCCC A GAGTCAGG	714	CCUGACUC CUGAUGAG X CGAA IGGAUUUU	3223
43	CCAGAGTC A GGGCCCTG	715	CAGGGCCC CUGAUGAG X CGAA IACUCUGG	3224
48	GTCAGGGC C CTGTACTT	716	AAGUACAG CUGAUGAG X CGAA ICCUCUGAC	3225
49	TCAGGGCC C TGTACTTT	717	AAAGUACA CUGAUGAG X CGAA IGCCUGA	3226
50	CAGGGCCC T GTACTTTC	718	GAAAGUAC CUGAUGAG X CGAA IGCCUCUG	3227
55	CCCTGTAC T TTCCTGCT	719	AGCAGGAA CUGAUGAG X CGAA IUACAGGG	3228
59	GTAATTTC C TGCTGGTG	720	CACCAGCA CUGAUGAG X CGAA IAAAGUAC	3229
60	TACTTTCC T GCTGGTGG	721	CCACCAGC CUGAUGAG X CGAA IGAAAGUA	3230
63	TTTCCTGC T GGTGGCTC	722	GAGCCACC CUGAUGAG X CGAA ICAGGAAA	3231
70	CTGGTGGC T CCAGTTCA	723	UGAACUGG CUGAUGAG X CGAA ICCACCAG	3232
72	GGTGGCTC C AGTTCAGG	724	CCUGAACU CUGAUGAG X CGAA IAGCCACC	3233
73	GTGGCTCC A GTTCAGGA	725	UCCUGAAC CUGAUGAG X CGAA IGAGCCAC	3234
78	TCCAGTTC A GGAACAGT	726	ACUGUUC CUGAUGAG X CGAA IAACUGGA	3235
84	TCAGGAAC A GTGAGCCC	727	GGGUCAC CUGAUGAG X CGAA IUUCCUGA	3236
91	CAGTGAGC C TGCTCAG	728	CUGAGCAG CUGAUGAG X CGAA ICUCACUG	3237
92	AGTGAGCC C TGCTCAGA	729	UCUGAGCA CUGAUGAG X CGAA IGCUCACU	3238
93	GTGAGCCC T GCTCAGAA	730	UUCUGAGC CUGAUGAG X CGAA IGGCUCAC	3239
96	AGCCCTGC T CAGAATAC	731	GUAUUCUG CUGAUGAG X CGAA ICAGGGCU	3240
98	CCCTGCTC A GAATACTG	732	CAGUAUUC CUGAUGAG X CGAA IAGCAGGG	3241
105	CAGAATAC T GTCTCTGC	733	GCAGAGAC CUGAUGAG X CGAA IUUUCUG	3242
109	ATACTGTC T CTGCCATA	734	UAUGGCAG CUGAUGAG X CGAA IACAGUUA	3243
111	ACTGTCTC T GCCATATC	735	GAUAUGGC CUGAUGAG X CGAA IAGACAGU	3244
114	GTCTCTGC C ATATCGTC	736	GACGAUUA CUGAUGAG X CGAA ICAGAGAC	3245
115	TCTCTGCC A TATCGTCA	737	UGACGAUA CUGAUGAG X CGAA IGCAGAGA	3246
123	ATATCGTC A ATCTTATC	738	GAUAAGAU CUGAUGAG X CGAA IACGAUUA	3247
127	CGTCAATC T TATCGAAG	739	CUUCGAUA CUGAUGAG X CGAA IAUUGACG	3248
138	TCGAAGAC T GGGGACCC	740	GGGUCCCC CUGAUGAG X CGAA IUCUUCGA	3249
145	CTGGGGAC C CTGTACCG	741	CGGUACAG CUGAUGAG X CGAA IUCCCCAG	3250
146	TGGGGACC C TGTACCGA	742	UCGGUACA CUGAUGAG X CGAA IGUCCCCA	3251
147	GGGGACCC T GTACCGAA	743	UUCGGUAC CUGAUGAG X CGAA IGGUCCCC	3252
152	CCCTGTAC C GAACATGG	744	CCAUGUUC CUGAUGAG X CGAA IUACAGGG	3253
157	TACCGAAC A TGGAGAAC	745	GUUCUCCA CUGAUGAG X CGAA IUUCGGUA	3254
166	TGGAGAAC A TCGCATCA	746	UGAUGCGA CUGAUGAG X CGAA IUUCUCCA	3255
171	AACATCGC A TCAGGACT	747	AGUCCUGA CUGAUGAG X CGAA ICGAUGUU	3256
174	ATCGCATC A GGAATCCT	748	AGGAGUCC CUGAUGAG X CGAA IAUCCGAU	3257

Table 38

179	ATCAGGAC T CCTAGGAC	749	GUCCUAGG CUGAUGAG X CGAA IUCCUGAU	3258
181	CAGGACTC C TAGGACCC	750	GGGUCCUA CUGAUGAG X CGAA IAGUCCUG	3259
182	AGGACTCC T AGGACCCC	751	GGGGUCCU CUGAUGAG X CGAA IGAGUCCU	3260
188	CCTAGGAC C CCTGCTCG	752	CGAGCAGG CUGAUGAG X CGAA IUCCUAGG	3261
189	CTAGGACC C CTGCTCGT	753	ACGAGCAG CUGAUGAG X CGAA IGUCCUAG	3262
190	TAGGACCC C TGCTCGTG	754	CACGAGCA CUGAUGAG X CGAA IGGUCCUA	3263
191	AGGACCCC T GCTCGTGT	755	ACACGAGC CUGAUGAG X CGAA IGGGUCCU	3264
194	ACCCCTGC T CGTGTAC	756	GUAACACG CUGAUGAG X CGAA ICAGGGGU	3265
203	CGTGTAC A GGCGGGGT	757	ACCCCGCC CUGAUGAG X CGAA IUAACACG	3266
217	GGTTTTTC T TGTTGACA	758	UGUCAACA CUGAUGAG X CGAA IAAAAACC	3267
225	TTGTTGAC A AAAATCCT	759	AGGAUUUU CUGAUGAG X CGAA IUCAACAA	3268
232	CAAAAATC C TCACAATA	760	UAUUGUGA CUGAUGAG X CGAA IAUUUUUG	3269
233	AAAAATCC T CACAATAC	761	GUUUGUG CUGAUGAG X CGAA IGAUUUUU	3270
235	AAATCCTC A CAATACCA	762	UGGUAUUG CUGAUGAG X CGAA IAGGAUUU	3271
237	ATCCTCAC A ATACCACA	763	UGUGUAU CUGAUGAG X CGAA IUGAGGAU	3272
242	CACAATAC C ACAGAGTC	764	GACUCUGU CUGAUGAG X CGAA IUUUGUG	3273
243	ACAATACC A CAGAGTCT	765	AGACUCUG CUGAUGAG X CGAA IGUAUUGU	3274
245	AATACCAC A GAGTCTAG	766	CUAGACUC CUGAUGAG X CGAA IUGGUAUU	3275
251	ACAGAGTC T AGACTCGT	767	ACGAGUCU CUGAUGAG X CGAA IACUCUGU	3276
256	GTCTAGAC T CGTGGTGG	768	CCACCACG CUGAUGAG X CGAA IUCUAGAC	3277
267	TGGTGGAC T TCTCTCAA	769	UUGAGAGA CUGAUGAG X CGAA IUCCACCA	3278
270	TGGACTTC T CTCAATT	770	AAAUUGAG CUGAUGAG X CGAA IAAGUCCA	3279
272	GACTTCTC T CAATTTTC	771	GAAAAUUG CUGAUGAG X CGAA IAGAAGUC	3280
274	CTTCTCTC A ATTTTCTA	772	UAGAAAAU CUGAUGAG X CGAA IACAGAAG	3281
281	CAATTTTC T AGGGGGAA	773	UUCCCCCU CUGAUGAG X CGAA IAAAAUUG	3282
291	GGGGGAAC A CCCGTGTG	774	CACACGGG CUGAUGAG X CGAA IUUCCCCC	3283
293	GGGAACAC C CGTGTGTC	775	GACACACG CUGAUGAG X CGAA IUGUCCCC	3284
294	GGAACACC C GTGTGTCT	776	AGACACAC CUGAUGAG X CGAA IGUGUCCC	3285
302	CGTGTGTC T TGGCCAAA	777	UUUGGCCA CUGAUGAG X CGAA IACACACG	3286
307	GTCTTGGC C AAAATTCG	778	CGAAUUUU CUGAUGAG X CGAA ICCAAGAC	3287
308	TCTTGCC A AAATTCGC	779	GCGAAUUU CUGAUGAG X CGAA IGCCAAGA	3288
317	AAATTCGC A GTCCCAAA	780	UUUGGGAC CUGAUGAG X CGAA ICGAAUUU	3289
321	TCGCAGTC C CAAATCTC	781	GAGAUUUG CUGAUGAG X CGAA IACUGCGA	3290
322	CGCAGTCC C AAATCTCC	782	GGAGAUUU CUGAUGAG X CGAA IGACUGCG	3291
323	GCAGTCCC A AATCTCCA	783	UGGAGAUU CUGAUGAG X CGAA IGGACUGC	3292
328	CCCAAATC T CCAGTCAC	784	GUGACUGG CUGAUGAG X CGAA IAUUUGGG	3293
330	CAAATCTC C AGTCACTC	785	GAGUGACU CUGAUGAG X CGAA IAGAUUUG	3294
331	AAATCTCC A GTCACCTA	786	UGAGUGAC CUGAUGAG X CGAA IGAGAUUU	3295
335	CTCCAGTC A CTCACCAA	787	UUGGUGAG CUGAUGAG X CGAA IACUGGAG	3296
337	CCAGTCAC T CACCAACC	788	GGUUGGUG CUGAUGAG X CGAA IUGACUGG	3297
339	AGTCACTC A CCAACCTG	789	CAGGUUGG CUGAUGAG X CGAA IAGUGACU	3298
341	TCACTCAC C AACCTGTT	790	AACAGGUU CUGAUGAG X CGAA IUAGUGA	3299
342	CACTCAC C AACTGTTG	791	CAACAGGU CUGAUGAG X CGAA IGUGAGUG	3300
345	TCACCAAC C TGTGTGCC	792	GGACAACA CUGAUGAG X CGAA IUUGGUGA	3301
346	CACCAACC T GTGTCTCT	793	AGGACAAC CUGAUGAG X CGAA IGUGGUG	3302
353	CTGTTGTC C TCCAATTT	794	AAAUUGGA CUGAUGAG X CGAA IACAACAG	3303
354	TGTTGTCC T CCAATTTG	795	CAAAUUGG CUGAUGAG X CGAA IGACAACA	3304
356	TTGTCCTC C AATTTGTC	796	GACAAAUU CUGAUGAG X CGAA IAGGACAA	3305
357	TGTCCTCC A ATTTGTCC	797	GGACAAAU CUGAUGAG X CGAA IGAGGACA	3306
365	AATTTGTC C TGTTATC	798	GAUAACCA CUGAUGAG X CGAA IACAAAUU	3307
366	ATTTGTCC T GGTATCG	799	CGAUAACC CUGAUGAG X CGAA IGACAAAU	3308

Table 38

376	GTTATCGC T GGATGTGT	800	ACACAUC CUGAUGAG X CGAA ICGAUAAC	3309
386	GATGTGTC T GCGGCGTT	801	AACGCCGC CUGAUGAG X CGAA IACACAUC	3310
400	GTTTTATC A TCTTCCTC	802	GAGGAAGA CUGAUGAG X CGAA IAUAAAAC	3311
403	TTATCATC T TCCTCTGC	803	GCAGAGGA CUGAUGAG X CGAA IAUGAUAA	3312
406	TCATCTTC C TCTGCATC	804	GAUGCAGA CUGAUGAG X CGAA IAAGAUGA	3313
407	CATCTTCC T CTGCATCC	805	GGAUGCAG CUGAUGAG X CGAA IGAAGAUG	3314
409	TCTTCCTC T GCATCCTG	806	CAGGAUGC CUGAUGAG X CGAA IAGGAAGA	3315
412	TCTCTGCA A TCCTGCTG	807	CAGCAGGA CUGAUGAG X CGAA ICAGAGGA	3316
415	TCTGCATC C TGCTGCTA	808	UAGCAGCA CUGAUGAG X CGAA IAUGCAGA	3317
416	CTGCATCC T GCTGCTAT	809	AUAGCAGC CUGAUGAG X CGAA IGAUGCAG	3318
419	CATCCTGC T GCTATGCC	810	GGCAUAGC CUGAUGAG X CGAA ICAGGAUG	3319
422	CCTGCTGC T ATGCCTCA	811	UGAGGCAU CUGAUGAG X CGAA ICAGCAGG	3320
427	TGCTATGC C TCATCTTC	812	GAAGAUGA CUGAUGAG X CGAA ICAUAGCA	3321
428	GCTATGCC T CATCTTCT	813	AGAAGAUG CUGAUGAG X CGAA IGAUAGC	3322
430	TATGCCTC A TCTTCTTG	814	CAAGAAGA CUGAUGAG X CGAA IAGGCAUA	3323
433	GCCTCATC T TCTTGTTG	815	CAACAAGA CUGAUGAG X CGAA IAUGAGGC	3324
436	TCATCTTC T TGTTGGTT	816	AACCAACA CUGAUGAG X CGAA IAAGAUGA	3325
446	GTTGGTTC T TCTGGACT	817	AGUCCAGA CUGAUGAG X CGAA IAACCAAC	3326
449	GGTTCCTC T GGAATATC	818	GAUAGUCC CUGAUGAG X CGAA IAAGAACC	3327
454	TTCTGGAC T ATCAAGGT	819	ACCUUGAU CUGAUGAG X CGAA IUCCAGAA	3328
458	GGACTATC A AGGTATGT	820	ACAUACCU CUGAUGAG X CGAA IAUAGUCC	3329
470	TATGTTGC C CGTTTGTC	821	GACAAACG CUGAUGAG X CGAA ICAACAUA	3330
471	ATGTTGCC C GTTGTGCC	822	GGACAAAC CUGAUGAG X CGAA IGCAACAU	3331
479	CGTTTGTC C TCTAATTC	823	GAAUUGA CUGAUGAG X CGAA IACAAACG	3332
480	GTTTGTC C TCAATTC	824	GGAAUUG CUGAUGAG X CGAA IGACAAAC	3333
482	TTGTCTC T AATTCAG	825	CUGGAAU CUGAUGAG X CGAA IAGGACAA	3334
488	TCTAATTC C AGGATCAT	826	AUGAUCCU CUGAUGAG X CGAA IAAUUGA	3335
489	CTAATTC A GGATCATC	827	GAUGAUCC CUGAUGAG X CGAA IGAUUG	3336
495	CCAGGATC A TCAACAAC	828	GUUGUUGA CUGAUGAG X CGAA IAUCUGG	3337
498	GGATCATC A ACAACCAG	829	CUGGUUGU CUGAUGAG X CGAA IAUAGUCC	3338
501	TCATCAAC A ACCAGCAC	830	GUGGUGGU CUGAUGAG X CGAA IUUGAUGA	3339
504	TCAACAAC C AGCACCGG	831	CCGGUGCU CUGAUGAG X CGAA IUUGUUGA	3340
505	CAACAACC A GCACCGGA	832	UCCGGUGC CUGAUGAG X CGAA IGUUGUG	3341
508	CAACCAGC A CCGACCA	833	UGGUCCGG CUGAUGAG X CGAA ICUGGUUG	3342
510	ACCAGCAC C GGACCATG	834	CAUGGUCC CUGAUGAG X CGAA IUGGUGU	3343
515	CACCGGAC C ATGCAAAA	835	UUUUGCAU CUGAUGAG X CGAA IUCCGGUG	3344
516	ACCGGACC A TGCAAAAC	836	GUUUGCA CUGAUGAG X CGAA IGUCCGGU	3345
520	GACCATGC A AAACCTGC	837	GCAGGUUU CUGAUGAG X CGAA ICAUGGUC	3346
525	TGCAAAAC C TGACAAC	838	GUUGUGCA CUGAUGAG X CGAA IUUUUGCA	3347
526	GCAAAACC T GCACAACT	839	AGUUGUGC CUGAUGAG X CGAA IGUUUUGC	3348
529	AAACCTGC A CAACTCCT	840	AGGAGUUG CUGAUGAG X CGAA ICAGGUUU	3349
531	ACCTGCAC A ACTCCTGC	841	GCAGGAGU CUGAUGAG X CGAA IUGCAGGU	3350
534	TGCACAAC T CCTGCTCA	842	UGAGCAGG CUGAUGAG X CGAA IUUGUGCA	3351
536	CACAACCT C TGCTCAAG	843	CUUGAGCA CUGAUGAG X CGAA IAGUUGUG	3352
537	ACAACCTC T GCTCAAGG	844	CCUUGAGC CUGAUGAG X CGAA IGAGUUGU	3353
540	ACTCCTGC T CAAGGAAC	845	GUUCCUUG CUGAUGAG X CGAA ICAGGAGU	3354
542	TCCTGCTC A AGGAACCT	846	AGGUUCCU CUGAUGAG X CGAA IAGCAGGA	3355
549	CAAGGAAC C TCTATGTT	847	AACAUAGA CUGAUGAG X CGAA IUUCCUUG	3356
550	AAGGAACC T CTATGTTT	848	AAACAUG CUGAUGAG X CGAA IGUCCUUG	3357
552	GGAACCTC T ATGTTTCC	849	GGAAACAU CUGAUGAG X CGAA IAGGUUCC	3358
560	TATGTTTC C CTCATGTT	850	AACAUGAG CUGAUGAG X CGAA IAAACAUA	3359

Table 38

561	ATGTTTCC C TCATGTTG	851	CAACAUGA CUGAUGAG X CGAA IGAAACAU	3360
562	TGTTTCCC T CATGTTGC	852	GCAACAUG CUGAUGAG X CGAA IGGAAACA	3361
564	TTTCCCTC A TGTGCTG	853	CAGCAACA CUGAUGAG X CGAA IAGGGAAA	3362
571	CATGTTGC T GTACAAAA	854	UUUUGUAC CUGAUGAG X CGAA ICAACAUG	3363
576	TGCTGTAC A AAACCTAC	855	GUAGGUUU CUGAUGAG X CGAA IUACAGCA	3364
581	TACAAAAC C TACGGACG	856	CGUCCGUA CUGAUGAG X CGAA IUUUUGUA	3365
582	ACAAAACC T ACGGACGG	857	CCGUCCGU CUGAUGAG X CGAA IGUUUUGU	3366
595	ACGGAAAC T GCACCTGT	858	ACAGGUGC CUGAUGAG X CGAA IUUCCGU	3367
598	GAAACTGC A CCTGTATT	859	AAUACAGG CUGAUGAG X CGAA ICAGUUC	3368
600	AACTGCAC C TGTATTCC	860	GGAAUACA CUGAUGAG X CGAA IUGCAGUU	3369
601	ACTGCACC T GTATTCCC	861	GGGAAUAC CUGAUGAG X CGAA IGUGCAGU	3370
608	CTGTATT C CATCCCAT	862	AUGGGAUG CUGAUGAG X CGAA IAAUACAG	3371
609	TGTATTCC C ATCCATC	863	GAUGGGAU CUGAUGAG X CGAA IGAUACA	3372
610	GTATTCCC A TCCCATCA	864	UGAUGGGA CUGAUGAG X CGAA IGGAAUAC	3373
613	TTCCCATC C CATCATCT	865	AGAUGAUG CUGAUGAG X CGAA IAUGGGAA	3374
614	TCCCATCC C ATCATCTT	866	AAGAUGAU CUGAUGAG X CGAA IGAUGGGA	3375
615	CCCATCCC A TCATCTTG	867	CAAGAUGA CUGAUGAG X CGAA IGGAUGGG	3376
618	ATCCCATC A TCTTGGGC	868	GCCCAAGA CUGAUGAG X CGAA IAUGGGAU	3377
621	CCATCATC T TGGGCTTT	869	AAAGCCCA CUGAUGAG X CGAA IAUGAUGG	3378
627	TCTTGGGC T TTCGAAA	870	UUUGCGAA CUGAUGAG X CGAA ICCCAAGA	3379
633	GCTTTCGC A AAATACCT	871	AGGUUUU CUGAUGAG X CGAA ICGAAAGC	3380
640	CAAATAC C TATGGGAG	872	CUCCCAUA CUGAUGAG X CGAA IUUUUUUG	3381
641	AAAATACC T ATGGGAGT	873	ACUCCCAU CUGAUGAG X CGAA IGUAUUUU	3382
654	GAGTGGGC C TCAGTCCG	874	CGGACUGA CUGAUGAG X CGAA ICCCACUC	3383
655	AGTGGGCC T CAGTCCGT	875	ACGGACUG CUGAUGAG X CGAA IGCCCACU	3384
657	TGGGCCTC A GTCCGTTT	876	AAACGGAC CUGAUGAG X CGAA IAGGCCCA	3385
661	CCTCAGTC C GTTCTCT	877	AGAGAAAC CUGAUGAG X CGAA IACUGAGG	3386
667	TCCGTTTC T CTTGGCTC	878	GAGCCAAG CUGAUGAG X CGAA IAAACGGA	3387
669	CGTTTCTC T TGGCTCAG	879	CUGAGCCA CUGAUGAG X CGAA IAGAAACG	3388
674	CTCTTGGC T CAGTTTAC	880	GUAAACUG CUGAUGAG X CGAA ICCAAGAG	3389
676	CTTGCTC A GTTACTA	881	UAGUAAAC CUGAUGAG X CGAA IAGCCAAG	3390
683	CAGTTTAC T AGTGCCAT	882	AUGGCACU CUGAUGAG X CGAA IUAAACUG	3391
689	ACTAGTGC C ATTTGTTT	883	GAACAAAU CUGAUGAG X CGAA ICACUAGU	3392
690	CTAGTGCC A TTTGTTCA	884	UGAACAAA CUGAUGAG X CGAA IGCACUAG	3393
698	ATTTGTTT A GTGGTTTC	885	CGAACCAC CUGAUGAG X CGAA IAACAAAU	3394
713	CGTAGGGC T TTCCCCCA	886	UGGGGGAA CUGAUGAG X CGAA ICCCUACG	3395
717	GGGCTTTC C CCCACTGT	887	ACAGUGGG CUGAUGAG X CGAA IAAAGCCC	3396
718	GGCTTTCC C CCACTGTC	888	GACAGUGG CUGAUGAG X CGAA IGAAAGCC	3397
719	GCTTTCCC C CACTGTCT	889	AGACAGUG CUGAUGAG X CGAA IGGAAAGC	3398
720	CTTTCCCC C ACTGTCTG	890	CAGACAGU CUGAUGAG X CGAA IGGGAAAG	3399
721	TTTCCCCC A CTGTCTGG	891	CCAGACAG CUGAUGAG X CGAA IGGGGAAA	3400
723	TCCCCCAC T GTCTGGCT	892	AGCCAGAC CUGAUGAG X CGAA IUUGGGGA	3401
727	CCACTGTC T GGCTTTCA	893	UGAAAGCC CUGAUGAG X CGAA IACAGUGG	3402
731	TGTCTGGC T TTCAGTTA	894	UAACUGAA CUGAUGAG X CGAA ICCAGACA	3403
735	TGGCTTTC A GTTATATG	895	CAUAUAAC CUGAUGAG X CGAA IAAAGCCA	3404
764	TTGGGGGC C AAGTCTGT	896	ACAGACUU CUGAUGAG X CGAA ICCCCCAA	3405
765	TGGGGGCC A AGTCTGTA	897	UACAGACU CUGAUGAG X CGAA IGCCCCCA	3406
770	GCCAAGTC T GTACAACA	898	UGUUGUAC CUGAUGAG X CGAA IACUUGGC	3407
775	GTCTGTAC A ACATCTTG	899	CAAGAUGU CUGAUGAG X CGAA IUACAGAC	3408
778	TGTACAAC A TCTTGAGT	900	ACUCAAGA CUGAUGAG X CGAA IUUGUACA	3409
781	ACAACATC T TGAGTCCC	901	GGGACUCA CUGAUGAG X CGAA IAUGUUGU	3410

Table 38

788	CTTGAGTC C CTTTATGC	902	GCAUAAAG CUGAUGAG X CGAA IACUCAAG	3411
789	TTGAGTCC C TTTATGCC	903	GGCAUAAA CUGAUGAG X CGAA IGACUCA	3412
790	TGAGTCCC T TTATGCCG	904	CGGCAUAA CUGAUGAG X CGAA IGGACUCA	3413
797	CTTTATGC C GCTGTTAC	905	GUAACAGC CUGAUGAG X CGAA ICAUAAAG	3414
800	TATGCCGC T GTTACCAA	906	UUGGUAAAC CUGAUGAG X CGAA ICGGCAUA	3415
806	GCTGTTAC C AATTTTCT	907	AGAAAAUU CUGAUGAG X CGAA IUAACAGC	3416
807	CTGTTACC A ATTTTCTT	908	AAGAAAAU CUGAUGAG X CGAA IGUAACAG	3417
814	CAATTTTC T TTTGTCTT	909	AAGACAAA CUGAUGAG X CGAA IAAAAUUG	3418
821	CTTTTGTC T TTGGGTAT	910	AUACCCAA CUGAUGAG X CGAA IACAAAAG	3419
832	GGGTATAC A TTAAACC	911	GGUUUAAA CUGAUGAG X CGAA IUUAACCC	3420
840	ATTTAAAC C CTCACAAA	912	UUUGUGAG CUGAUGAG X CGAA IUUUAAA	3421
841	TTTAAACC C TCACAAA	913	UUUUGUGA CUGAUGAG X CGAA IGUUUAAA	3422
842	TTAAACCC T CACAAAAC	914	GUUUUGUG CUGAUGAG X CGAA IGGUUUAA	3423
844	AAACCCTC A CAAAACAA	915	UUGUUUUG CUGAUGAG X CGAA IAGGGUUU	3424
846	ACCCTCAC A AAACAAA	916	UUUUGUUU CUGAUGAG X CGAA IUGAGGGU	3425
851	CACAAAAC A AAAAGATG	917	CAUCUUUU CUGAUGAG X CGAA IUUUUGUG	3426
869	GGATATTC C CTTAACTT	918	AAGUUAAG CUGAUGAG X CGAA IAAUAUCC	3427
870	GATATTC C TTAACCTC	919	GAAGUUA CUGAUGAG X CGAA IGAAUAUC	3428
871	ATATTC C TAACTTCA	920	UGAAGUUA CUGAUGAG X CGAA IGGAAUAU	3429
876	CCCTTAAC T TCATGGGA	921	UCCCAUGA CUGAUGAG X CGAA IUUAAGGG	3430
879	TTAACTTC A TGGGATAT	922	AUAUCCCA CUGAUGAG X CGAA IAAGUUA	3431
906	GTTGGGGC A CATTGCCA	923	UGGCAUG CUGAUGAG X CGAA ICCCCAAC	3432
908	TGGGGCAC A TTGCCACA	924	UGUGGCAA CUGAUGAG X CGAA IUGCCCCA	3433
913	CACATTGC C ACAGGAAC	925	GUUCCUGU CUGAUGAG X CGAA ICAAUGUG	3434
914	ACATTGCC A CAGGAACA	926	UGUCCUG CUGAUGAG X CGAA IUGGCAAU	3435
916	ATTGCCAC A GGAACATA	927	UAUGUCC CUGAUGAG X CGAA IUUGCAAU	3436
922	ACAGGAAC A TATTGTAC	928	GUACAAUA CUGAUGAG X CGAA IUUCCUGU	3437
931	TATTGTAC A AAAAATCA	929	UGAUUUUU CUGAUGAG X CGAA IUACAAUA	3438
939	AAAAAATC A AAATGTGT	930	ACACAUUU CUGAUGAG X CGAA IAUUUUU	3439
958	TAGGAAAC T TCCTGTAA	931	UUAACAGGA CUGAUGAG X CGAA IUUCCUA	3440
961	GAAACTTC C TGTAACAC	932	UGUUUACA CUGAUGAG X CGAA IAAGUUUC	3441
962	AAACTTCC T GTAACACAG	933	CUGUUUAC CUGAUGAG X CGAA IGAAGUUU	3442
969	CTGTAAAC A GGCCTATT	934	AAUAGGCC CUGAUGAG X CGAA IUUUACAG	3443
973	AAACAGGC C TATTGATT	935	AAUCAUA CUGAUGAG X CGAA ICCUGUUU	3444
974	AACAGGCC T ATTGATTG	936	CAAUCAAU CUGAUGAG X CGAA IGCCUGUU	3445
994	AGTATGTC A ACGAATTG	937	CAAUUCGU CUGAUGAG X CGAA IACUAUCU	3446
1009	TGTGGGTC T TTTGGGGT	938	ACCCCAA CUGAUGAG X CGAA IACCCACA	3447
1022	GGGTTTGC C GCCCCTTT	939	AAAGGGGC CUGAUGAG X CGAA ICAAACCC	3448
1025	TTTGCCGC C CCTTTCAC	940	GUGAAAGG CUGAUGAG X CGAA ICGGCAA	3449
1026	TTGCCGCC C CTTTCACG	941	CGUGAAAG CUGAUGAG X CGAA IGGCGCA	3450
1027	TGCCGCC C TTTACGCA	942	GCGUGAAA CUGAUGAG X CGAA IGGCGGC	3451
1028	GCCGCCCC T TTTACGCA	943	UGCGUGAA CUGAUGAG X CGAA IGGCGGC	3452
1032	CCCCTTTC A CGCAATGT	944	ACAUUGCG CUGAUGAG X CGAA IAAAGGGG	3453
1036	TTTCACGC A ATGTGGAT	945	AUCCACAU CUGAUGAG X CGAA ICGUGAAA	3454
1049	GGATATTC T GCTTAAT	946	AUUAAAGC CUGAUGAG X CGAA IAAUAUCC	3455
1052	TATTCTGC T TTAATGCC	947	GGCAUUA CUGAUGAG X CGAA ICAGAAUA	3456
1060	TTTAATGC C TTTATATG	948	CAUAUAAA CUGAUGAG X CGAA ICAUUAUA	3457
1061	TTAATGCC T TTATATGC	949	GCAUAUAA CUGAUGAG X CGAA IGCAUUA	3458
1070	TTATATGC A TGCAACA	950	UGUAUGCA CUGAUGAG X CGAA ICAUAUA	3459
1074	ATGCATGC A TACAAGCA	951	UGCUGUA CUGAUGAG X CGAA ICAUGCAU	3460
1078	ATGCATAC A AGCAAAAC	952	GUUUUGCU CUGAUGAG X CGAA IUAUGCAU	3461

Table 38

1082	ATACAAGC A AAACAGGC	953	GCCUGUUU CUGAUGAG X CGAA ICUUGUAU	3462
1087	AGCAAAAC A GGCTTTTA	954	UAAAAGCC CUGAUGAG X CGAA IUUUUGCU	3463
1091	AAACAGGC T TTTACTTT	955	AAAGUAAA CUGAUGAG X CGAA ICCUGUUU	3464
1097	GCTTTTAC T TTCTCGCC	956	GGCGAGAA CUGAUGAG X CGAA IUAAAAGC	3465
1101	TTACTTTC T CGCCAAC	957	AGUUGGCG CUGAUGAG X CGAA IAAAGUAA	3466
1105	TTCTCGC C AACTTACA	958	UGUAAGUU CUGAUGAG X CGAA ICGAGAAA	3467
1106	TTCTCGCC A ACTTACAA	959	UUGUAAGU CUGAUGAG X CGAA ICGGAGAA	3468
1109	TCGCCAAC T TACAAGGC	960	GCCUUGUA CUGAUGAG X CGAA IUUGGCGA	3469
1113	CAACTTAC A AGGCCTTT	961	AAAGCCCU CUGAUGAG X CGAA IUAAGUUG	3470
1118	TACAAGGC C TTTCTAAG	962	CUUAGAAA CUGAUGAG X CGAA ICCUUGUA	3471
1119	ACAAGGCC T TTCTAAGT	963	ACUUGAAA CUGAUGAG X CGAA IGCCUUGU	3472
1123	GGCCTTTT T AAGTAAAC	964	GUUUACUU CUGAUGAG X CGAA IAAAGGCC	3473
1132	AAGTAAAC A GTATGTGA	965	UCACAUAC CUGAUGAG X CGAA IUUUACUU	3474
1143	ATGTGAAC C TTTACCCC	966	GGGGUAAA CUGAUGAG X CGAA IUUCACAU	3475
1144	TGTGAACC T TTACCCCG	967	CGGGGUAA CUGAUGAG X CGAA IGUUCACA	3476
1149	ACCTTTAC C CCGTTGCT	968	AGCAACGG CUGAUGAG X CGAA IUAAAGGU	3477
1150	CCTTTACC C CGTTGCTC	969	GAGCAACG CUGAUGAG X CGAA IGUAAAGG	3478
1151	CTTTACCC C GTTGCTCG	970	CGAGCAAC CUGAUGAG X CGAA IGGUAAAAG	3479
1157	CCCGTTGC T CGGCAACG	971	CGUUGCCG CUGAUGAG X CGAA ICAACGGG	3480
1162	TGCTCGGC A ACGGCCTG	972	CAGGCCGU CUGAUGAG X CGAA ICCGAGCA	3481
1168	GCAACGGC C TGGTCTAT	973	AUAGACCA CUGAUGAG X CGAA ICCGUUGC	3482
1169	CAACGGCC T GGTCTATG	974	CAUAGACC CUGAUGAG X CGAA IGCCGUUG	3483
1174	GCCTGGTC T ATGCCAAG	975	CUUGGCAU CUGAUGAG X CGAA IACCAGGC	3484
1179	GTCTATGC C AAGTGTTC	976	AAACACUU CUGAUGAG X CGAA ICAUAGAC	3485
1180	TCTATGCC A AGTGTTCG	977	CAAACUCU CUGAUGAG X CGAA IGCAUAGA	3486
1190	GTGTTTGC T GAGCAAC	978	GUUGCGUC CUGAUGAG X CGAA ICAAACAC	3487
1196	GCTGACGC A ACCCCCAC	979	GUGGGGGU CUGAUGAG X CGAA ICGUCAGC	3488
1199	GACGCAAC C CCCACTGG	980	CCAGUGGG CUGAUGAG X CGAA IUUGCGUC	3489
1200	ACGCAACC C CCACTGGT	981	ACCAGUGG CUGAUGAG X CGAA IGUUGCGU	3490
1201	CGCAACCC C CACTGGTT	982	AACCAGUG CUGAUGAG X CGAA IGGUUGCG	3491
1202	GCAACCCC C ACTGGTTG	983	CAACCAGU CUGAUGAG X CGAA IGGGUUGC	3492
1203	CAACCCC A CTGGTTGG	984	CCAACCAG CUGAUGAG X CGAA IGGGUUG	3493
1205	ACCCCAC T GGTGGGG	985	CCCCAACC CUGAUGAG X CGAA IUUGGGGU	3494
1215	GTTGGGGC T TGGCCATA	986	UAUGGCCA CUGAUGAG X CGAA ICCCCAAC	3495
1220	GGCTTGGC C ATAGGCCA	987	UGGCCUAU CUGAUGAG X CGAA ICCAAGCC	3496
1221	GCTTGGCC A TAGGCCAT	988	AUGGCCUA CUGAUGAG X CGAA IGCCAAGC	3497
1227	CCATAGGC C ATCAGCGC	989	GCGCUGAU CUGAUGAG X CGAA ICCUAUGG	3498
1228	CATAGGCC A TCAGCGCA	990	UGCGUGA CUGAUGAG X CGAA IGCCUAUG	3499
1231	AGGCCATC A GCGCATGC	991	GCAUGGCG CUGAUGAG X CGAA IAUGCCCU	3500
1236	ATCAGCGC A TGCGTGGA	992	UCCACGCA CUGAUGAG X CGAA ICGUGAU	3501
1247	CGTGGAAC C TTTGTGTC	993	GACACAAA CUGAUGAG X CGAA IUUCCACG	3502
1248	GTGGAACC T TTTGTGTC	994	AGACACAA CUGAUGAG X CGAA IGUUCCAC	3503
1256	TTTGTGTC T CCTCTGCC	995	GGCAGAGG CUGAUGAG X CGAA IACACAAA	3504
1258	TGTGTCTC C TCTGCCGA	996	UCGGCAGA CUGAUGAG X CGAA IAGACACA	3505
1259	GTGTCTCC T CTGCCGAT	997	AUCGGCAG CUGAUGAG X CGAA IGAGACAC	3506
1261	GTCTCTC T GCCGATCC	998	GGAUCGGC CUGAUGAG X CGAA IAGGAGAC	3507
1264	TCCTCTGC C GATCCATA	999	UAUGGAUC CUGAUGAG X CGAA ICAGAGGA	3508
1269	TGCCGATC C ATACCGCG	1000	CGCGGUUU CUGAUGAG X CGAA IAUCGGCA	3509
1270	GCCGATCC A TACCGCGG	1001	CCGCGGUA CUGAUGAG X CGAA IGAUCGGC	3510
1274	ATCCATAC C GCGGAAC	1002	AGUUCGCG CUGAUGAG X CGAA IUAUGGAU	3511
1282	CGCGGAAC T CCTAGCCG	1003	CGGCUAGG CUGAUGAG X CGAA IUUCCGCG	3512

Table 38

1284	CGGAACTC C TAGCCGCT	1004	AGCGGCUA CUGAUGAG X CGAA IAGUCCG	3513
1285	GGAAGTCC T AGCCGCTT	1005	AAGCGGCU CUGAUGAG X CGAA IGAGUCC	3514
1289	CTCCTAGC C GCTTGTTC	1006	AAACAAGC CUGAUGAG X CGAA ICUAGGAG	3515
1292	CTAGCCGC T TGTTCGTC	1007	GCAAAACA CUGAUGAG X CGAA ICGGCUAG	3516
1301	TGTTTTGC T CGCAGCAG	1008	CUGCUGCG CUGAUGAG X CGAA ICAAAACA	3517
1305	TTGCTCGC A GCAGGTCT	1009	AGACCUGC CUGAUGAG X CGAA ICGAGCAA	3518
1308	CTCGCAGC A GGTCTGGG	1010	CCCAGACC CUGAUGAG X CGAA ICUGCGAG	3519
1313	AGCAGGTC T GGGGCAAA	1011	UUUGCCCC CUGAUGAG X CGAA IACCUGCU	3520
1319	TCTGGGGC A AAATCAT	1012	AUGAGUUU CUGAUGAG X CGAA ICCCCAGA	3521
1324	GGCAAAAC T CATCGGGA	1013	UCCCGAUG CUGAUGAG X CGAA IUUUUGCC	3522
1326	CAAAACTC A TCGGGAAT	1014	AGUCCCGA CUGAUGAG X CGAA IAGUUUUG	3523
1334	ATCGGGAC T GACAATTC	1015	GAAUUGUC CUGAUGAG X CGAA IUCCCGAU	3524
1338	GGACTGAC A ATTCTGTC	1016	GACAGAAU CUGAUGAG X CGAA IUCAGUCC	3525
1343	GACAATTC T GTCGTGCT	1017	AGCACGAC CUGAUGAG X CGAA IAAUUGUC	3526
1351	TGTCGTGC T CTCCCGCA	1018	UGC GGAG CUGAUGAG X CGAA ICACGACA	3527
1353	TCGTGCTC T CCCGCAAA	1019	UUUGCGGG CUGAUGAG X CGAA IAGCACGA	3528
1355	GTGCTCTC C CGCAAATA	1020	UAUUUGCG CUGAUGAG X CGAA IAGAGCAC	3529
1356	TGCTCTCC C GCAAATAT	1021	AUAUUUGC CUGAUGAG X CGAA IGAGAGCA	3530
1359	TCTCCCGC A AATATACA	1022	UGUAUAUU CUGAUGAG X CGAA ICGGGAGA	3531
1367	AAATATAC A TCATTTC	1023	GGAAUAUG CUGAUGAG X CGAA IUUAUAUU	3532
1370	TATACATC A TTTCCATG	1024	CAUGGAAA CUGAUGAG X CGAA IAUGUAUA	3533
1375	ATCATTTT C ATGGCTGC	1025	GCAGCCAU CUGAUGAG X CGAA IAAAUGAU	3534
1376	TCATTTC A TGGCTGCT	1026	AGCAGCCA CUGAUGAG X CGAA IGAAAUGA	3535
1381	TCCATGGC T GCTAGGCT	1027	AGCCUAGC CUGAUGAG X CGAA ICCAUGGA	3536
1384	ATGCTGTC T AGGCTGTG	1028	CACAGCCU CUGAUGAG X CGAA ICAGCCAU	3537
1389	TGCTAGGC T GTGCTGCC	1029	GGCAGCAC CUGAUGAG X CGAA ICCUAGCA	3538
1394	GGCTGTGC T GCCAATCG	1030	CAGUUGGC CUGAUGAG X CGAA ICACAGCC	3539
1397	TGTGCTGC C AACTGGAT	1031	AUCCAGUU CUGAUGAG X CGAA ICAGCACA	3540
1398	GTGCTGCC A ACTGGATC	1032	GAUCCAGU CUGAUGAG X CGAA IGCAGCAC	3541
1401	CTGCCAAC T GGATCCTA	1033	UAGGAUCC CUGAUGAG X CGAA IUUGGCAG	3542
1407	ACTGGATC C TACGCGGG	1034	CCCGCGUA CUGAUGAG X CGAA IAUCCAGU	3543
1408	CTGGATCC T ACGCGGGA	1035	UCCCGCGU CUGAUGAG X CGAA IGAUCCAG	3544
1421	GGGACGTC C TTTGTTTA	1036	UAAACAAA CUGAUGAG X CGAA IACGUCCC	3545
1422	GGACGTCC T TTGTTTAC	1037	GUAAACAA CUGAUGAG X CGAA IGACGUCC	3546
1434	TTTACGTC C CGTCGGCG	1038	CGCCGACG CUGAUGAG X CGAA IACGUAAA	3547
1435	TTACGTCC C GTCGGCGC	1039	GCGCCGAC CUGAUGAG X CGAA IGACGUAA	3548
1444	GTCGGCGC T GAATCCCG	1040	CGGGAUUC CUGAUGAG X CGAA ICGCCGAC	3549
1450	GCTGAATC C CGCGGACG	1041	CGUCCGCG CUGAUGAG X CGAA IAUUCAGC	3550
1451	CTGAATCC C CGCGACGA	1042	UCGUCCGC CUGAUGAG X CGAA IGAUUCAG	3551
1461	CGGACGAC C CCTCCCGG	1043	CCGGGAGG CUGAUGAG X CGAA IUCGUCCG	3552
1462	GGACGACC C CTCCCGGG	1044	CCCGGGAG CUGAUGAG X CGAA IGUCGUCC	3553
1463	GACGACCC C TCCCGGGG	1045	CCCCGGGA CUGAUGAG X CGAA IGGUCGUC	3554
1464	ACGACCCC T CCCGGGGC	1046	GCCCCGGG CUGAUGAG X CGAA IGGGUCGU	3555
1466	GACCCCTC C CGGGGCCG	1047	CGGCCCCG CUGAUGAG X CGAA IAGGGGUC	3556
1467	ACCCCTCC C GGGGCCGC	1048	GCGGCCCC CUGAUGAG X CGAA IGAGGGGU	3557
1473	CCCGGGGC C GCTTGGGG	1049	CCCAAGC CUGAUGAG X CGAA ICCCGGGG	3558
1476	GGGGCCGC T TGGGGCTC	1050	GAGCCCCA CUGAUGAG X CGAA ICGGCCCC	3559
1483	CTTGGGGC T CTACCGCC	1051	GGCGGUAG CUGAUGAG X CGAA ICCCAAG	3560
1485	TGGGGCTC T ACCGCCCC	1052	CGGGCGGU CUGAUGAG X CGAA IAGCCCCA	3561
1488	GGCTCTAC C GCCCGCTT	1053	AAGCGGGC CUGAUGAG X CGAA IUAGAGCC	3562
1491	TCTACCGC C CGCTTCTC	1054	GAGAAGCG CUGAUGAG X CGAA ICGGUAGA	3563

Table 38

1492	CTACCGCC C GCTTCTCC	1055	GGAGAAGC CUGAUGAG X CGAA ICGGUAG	3564
1495	CCGCCCGC T TCTCCGCC	1056	GGCGGAGA CUGAUGAG X CGAA ICGGCGG	3565
1498	CCCGCTTC T CCGCCTAT	1057	AUAGGCGG CUGAUGAG X CGAA IAAGCGG	3566
1500	CGCTTCTC C GCCTATTG	1058	CAAUAGGC CUGAUGAG X CGAA IAGAAGCG	3567
1503	TTCTCCGC C TATTGTAC	1059	GUACAAUA CUGAUGAG X CGAA ICGGAGAA	3568
1504	TCTCCGCC T ATTGTACC	1060	GGUACAAU CUGAUGAG X CGAA ICGGAGA	3569
1512	TATTGTAC C GACCGTCC	1061	GGACGGUC CUGAUGAG X CGAA IUACAAUA	3570
1516	GTACCGAC C GTCCACGG	1062	CCGUGGAC CUGAUGAG X CGAA IUCGGUAC	3571
1520	CGACCGTC C ACGGGGCG	1063	CGCCCCGU CUGAUGAG X CGAA IACGGUCG	3572
1521	GACCGTCC A CGGGGGCG	1064	GCGCCCCG CUGAUGAG X CGAA IGACGGUC	3573
1530	CGGGGCGC A CCTCTCTT	1065	AAGAGAGG CUGAUGAG X CGAA ICGCCCCG	3574
1532	GGGCGCAC C TCTCTTTA	1066	UAAAGAGA CUGAUGAG X CGAA IUGCGCCC	3575
1533	GGCGCACC T CTCTTTAC	1067	GUAAAGAG CUGAUGAG X CGAA IGUGCGCC	3576
1535	CGCACCTC T CTTTACGC	1068	GCGUAAAG CUGAUGAG X CGAA IAGGUGCG	3577
1537	CACCTCTC T TTACGCGG	1069	CCGCGUAA CUGAUGAG X CGAA IAGAGGUG	3578
1548	ACGCGGAC T CCGCTCT	1070	AGACGGGG CUGAUGAG X CGAA IUCCGCGU	3579
1550	GCGGACTC C CCGTCTGT	1071	ACAGACGG CUGAUGAG X CGAA IAGUCCGC	3580
1551	CGGACTCC C CGTCTGTG	1072	CACAGACG CUGAUGAG X CGAA IGAGUCCG	3581
1552	GGACTCCC C GTCTGTGC	1073	GCACAGAC CUGAUGAG X CGAA IGGAGUCC	3582
1556	TCCCCGTC T GTGCC TTC	1074	GAAGGCAC CUGAUGAG X CGAA IACGGGGA	3583
1561	GTCTGTGC C TTCTCATC	1075	GAUGAGAA CUGAUGAG X CGAA ICACAGAC	3584
1562	TCTGTGCC T TCTCATCT	1076	AGAUGAGA CUGAUGAG X CGAA IGCACAGA	3585
1565	GTGCC TTC T CATCTGCC	1077	GGCAG AUG CUGAUGAG X CGAA IAAGGCAC	3586
1567	GCCTTCTC A TCTGCCG	1078	CCGGCAGA CUGAUGAG X CGAA IAGAAGGC	3587
1570	TTCTCATC T GCCGGACC	1079	GGUCCGGC CUGAUGAG X CGAA IAUGAGAA	3588
1573	TCATCTGC C GGACCGTG	1080	CACGGUCC CUGAUGAG X CGAA ICAGAGUA	3589
1578	TGCCGAC C GTGTGCAC	1081	GUGCACAC CUGAUGAG X CGAA IUCCGGCA	3590
1585	CGGTGTGC A CTTGCTT	1082	AAGCGAAG CUGAUGAG X CGAA ICACACGG	3591
1587	GTGTGCAC T TCGCTTCA	1083	UGAAGCGA CUGAUGAG X CGAA IUGCACAC	3592
1592	CACTTCGC T TCACCTCT	1084	AGAGGUGA CUGAUGAG X CGAA ICGAAGUG	3593
1595	TTCGCTTC A CCTCTGCA	1085	UGCAGAGG CUGAUGAG X CGAA IAAGCGAA	3594
1597	CGCTTCAC C TCTGCACG	1086	CGUGCAGA CUGAUGAG X CGAA IUGAAGCG	3595
1598	GCTTCACC T CTGCACGT	1087	ACGUGCAG CUGAUGAG X CGAA IGUGAAGC	3596
1600	TTCACCTC T GCACGTCG	1088	CGACGUGC CUGAUGAG X CGAA IAGGUGAA	3597
1603	ACCTCTGC A CGTCGCAT	1089	AUGCGACG CUGAUGAG X CGAA ICAGAGGU	3598
1610	CACGTCGC A TGGAGACC	1090	GGUCUCCA CUGAUGAG X CGAA ICGACGUG	3599
1618	ATGGAGAC C ACCGTGAA	1091	UUCACGGU CUGAUGAG X CGAA IUCUCCAU	3600
1619	TGGAGACC A CCGTGAAC	1092	GUUCACGG CUGAUGAG X CGAA IGUCUCCA	3601
1621	GAGACCAC C GTGAACGC	1093	GCGUUCAC CUGAUGAG X CGAA IUGGUCUC	3602
1630	GTGAACGC C CACAGGAA	1094	UUCUGUG CUGAUGAG X CGAA ICGUUCAC	3603
1631	TGAACGCC C ACAGGAAC	1095	GUUCCUGU CUGAUGAG X CGAA ICGGUUCA	3604
1632	GAACGCC A CAGGAACC	1096	GGUCCUG CUGAUGAG X CGAA IGGCGUUC	3605
1634	ACGCCAC A GGAACCTG	1097	CAGGUUCC CUGAUGAG X CGAA IUGGGCGU	3606
1640	ACAGGAAC C TGCCCAAG	1098	CUUGGGCA CUGAUGAG X CGAA IUUCCUGU	3607
1641	CAGGAACC T GCCCAAGG	1099	CCUUGGGC CUGAUGAG X CGAA IGUCCUG	3608
1644	GAACCTGC C CAAGGTCT	1100	AGACCUUG CUGAUGAG X CGAA ICAGGUUC	3609
1645	AACCTGCC C AAGGTCTT	1101	AAGACCUU CUGAUGAG X CGAA IGCAGGUU	3610
1646	ACCTGCC A AGGTCTTG	1102	CAAGACCU CUGAUGAG X CGAA IGGCAGGU	3611
1652	CCAAGGTC T TGCATAAG	1103	CUUAUGCA CUGAUGAG X CGAA IACCUUGG	3612
1656	GGTCTGTC A TAAGAGGA	1104	UCCUCUUA CUGAUGAG X CGAA ICAAGACC	3613
1666	AAGAGGAC T CTTGGACT	1105	AGUCCAAG CUGAUGAG X CGAA IUCCUCUU	3614

Table 38

1668	GAGGACTC T TGGACTTT	1106	AAAGUCCA CUGAUGAG X CGAA IAGUCCUC	3615
1674	TCTTGGAC T TTCAGCAA	1107	UUGCUGAA CUGAUGAG X CGAA IUCCAAGA	3616
1678	GGACTTTC A GCAATGTC	1108	GACAUUGC CUGAUGAG X CGAA IAAAGUCC	3617
1681	CTTTCAGC A ATGTCAAC	1109	GUUGACAU CUGAUGAG X CGAA ICUGAAAG	3618
1687	GCAATGTC A ACGACCGA	1110	UCGGUCGU CUGAUGAG X CGAA IACAUUGC	3619
1693	TCAACGAC C GACCTTGA	1111	UCAAGGUC CUGAUGAG X CGAA IUCGUUGA	3620
1697	CGACCGAC C TTGAGGCA	1112	UGCCUCAA CUGAUGAG X CGAA IUCGGUCG	3621
1698	GACCGACC T TGAGGCAT	1113	AUGCCUCA CUGAUGAG X CGAA IGUCGGUC	3622
1705	CTTGAGGC A TACTTCAA	1114	UUGAAGUA CUGAUGAG X CGAA ICCUCAAG	3623
1709	AGGCATAC T TCAAAGAC	1115	GUCUUUGA CUGAUGAG X CGAA IUAUGCCU	3624
1712	CATACTTC A AAGACTGT	1116	ACAGUCUU CUGAUGAG X CGAA IAAGUAUG	3625
1718	TCAAAGAC T GTGTGTTT	1117	AAACACAC CUGAUGAG X CGAA IUCUUUGA	3626
1769	TAAAGGTC T TTGTACTA	1118	UAGUACAA CUGAUGAG X CGAA IACCUUUA	3627
1776	CTTTGTAC T AGGAGGCT	1119	AGCCUCCU CUGAUGAG X CGAA IUACAAAG	3628
1784	TAGGAGGC T GTAGGCAT	1120	AUGCCUAC CUGAUGAG X CGAA ICCUCCUA	3629
1791	CTGTAGGC A TAAATTGG	1121	CCAAUUUA CUGAUGAG X CGAA ICCUACAG	3630
1807	GTGTGTTC A CCAGCACC	1122	GGUGCUGG CUGAUGAG X CGAA IAACACAC	3631
1809	GTGTTCAC C AGCACCAT	1123	AUGGUGCU CUGAUGAG X CGAA IUGAACAC	3632
1810	TGTTCAAC A GCACCATG	1124	CAUGGUGC CUGAUGAG X CGAA IGUGAACA	3633
1813	TCACCAGC A CCATGCAA	1125	UUGCAUGG CUGAUGAG X CGAA ICUGGUGA	3634
1815	ACCAGCAC C ATGCAACT	1126	AGUUGCAU CUGAUGAG X CGAA IUGCUGGU	3635
1816	CCAGCACC A TGCAACTT	1127	AAGUUGCA CUGAUGAG X CGAA IGUCGUGG	3636
1820	CACCATGC A ACTTTTTC	1128	GAAAAAGU CUGAUGAG X CGAA ICAUGGUG	3637
1823	CATGCAAC T TTTTCACC	1129	GGUGAAAA CUGAUGAG X CGAA IUUGCAUG	3638
1829	ACTTTTTC A CCTCTGCC	1130	GGCAGAGG CUGAUGAG X CGAA IAAAAAGU	3639
1831	TTTTTCAC C TCTGCCTA	1131	UAGGCAGA CUGAUGAG X CGAA IUGAAAAA	3640
1832	TTTTCAAC T CTGCCTAA	1132	UUAGGCAG CUGAUGAG X CGAA IGUGAAAA	3641
1834	TTCAACCT C GCCTAATC	1133	GAUUAGGC CUGAUGAG X CGAA IAGGUGAA	3642
1837	ACCTCTGC C TAATCATC	1134	GAUGAUUA CUGAUGAG X CGAA ICAGAGGU	3643
1838	CCTCTGCC T AATCATCT	1135	AGAUGAUU CUGAUGAG X CGAA IGCAGAGG	3644
1843	GCCTAATC A TCTCATGT	1136	ACAUGAGA CUGAUGAG X CGAA IAUUAGGC	3645
1846	TAATCATC T CATGTTCA	1137	UGAACAUU CUGAUGAG X CGAA IAUUAUUA	3646
1848	ATCATCTC A TGTTCATG	1138	CAUGAACA CUGAUGAG X CGAA IAGAUGAU	3647
1854	TCATGTTT A TGTCCTAC	1139	GUAGGACA CUGAUGAG X CGAA IAACAUGA	3648
1859	TTCATGTC C TACTGTTT	1140	GAACAGUA CUGAUGAG X CGAA IACAUGAA	3649
1860	TCATGTCC T ACTGTTCA	1141	UGAACAGU CUGAUGAG X CGAA IGACAUGA	3650
1863	TGTCCTAC T GTTCAAGC	1142	GCUUGAAC CUGAUGAG X CGAA IUAGGACA	3651
1868	TACTGTTT A AGCCTCCA	1143	UGGAGGCU CUGAUGAG X CGAA IAACAGUA	3652
1872	GTTCAAGC C TCCAAGCT	1144	AGCUUGGA CUGAUGAG X CGAA ICUUGAAC	3653
1873	TTCAAGCC T CCAAGCTG	1145	CAGCUUGG CUGAUGAG X CGAA IGCUGGAA	3654
1875	CAAGCCTC C AAGCTGTG	1146	CACAGCUU CUGAUGAG X CGAA IAGGCUUG	3655
1876	AAGCCTCC A AGCTGTGC	1147	GCACAGCU CUGAUGAG X CGAA IGAGGCUU	3656
1880	CTCCAAGC T GTGCCTTG	1148	CAAGGCAC CUGAUGAG X CGAA ICUUGGAG	3657
1885	AGCTGTGC C TTGGGTGG	1149	CCACCCAA CUGAUGAG X CGAA ICACAGCU	3658
1886	GCTGTGCC T TGGGTGGC	1150	GCCACCCA CUGAUGAG X CGAA IGCACAGC	3659
1895	TGGGTGGC T TTGGGGCA	1151	UGCCCCAA CUGAUGAG X CGAA ICCACCCA	3660
1903	TTTGGGGC A TGGACATT	1152	AAUGUCCA CUGAUGAG X CGAA ICCCCAAA	3661
1909	GCATGGAC A TTGACCCG	1153	CGGGUCAA CUGAUGAG X CGAA IUCCAUGC	3662
1915	ACATTGAC C CGTATAAA	1154	UUUAUACG CUGAUGAG X CGAA IUCAAUGU	3663
1916	CATTGACC C GTATAAAG	1155	CUUUAUAC CUGAUGAG X CGAA IGUCAAUG	3664
1935	TTTGAGAC T TCTGTGGA	1156	UCCACAGA CUGAUGAG X CGAA ICUCCAAA	3665

Table 38

1938	GGAGCTTC T GTGGAGTT	1157	AACUCCAC CUGAUGAG X CGAA IAAGCUCC	3666
1949	GGAGTTAC T CTCTTTT	1158	AAAAAGAG CUGAUGAG X CGAA IUAACUCC	3667
1951	AGTTACTC T CTTTTTG	1159	CAAAAAAG CUGAUGAG X CGAA IAGUAACU	3668
1953	TTACTCTC T TTTTGGCC	1160	GGCAAAAA CUGAUGAG X CGAA IAGAGUAA	3669
1961	TTTTTTC C TTCTGACT	1161	AGUCAGAA CUGAUGAG X CGAA ICAAAAA	3670
1962	TTTTTGCC T TCTGACTT	1162	AAGUCAGA CUGAUGAG X CGAA IGCAAAAA	3671
1965	TTGCCCTC T GACTTCTT	1163	AAGAAGUC CUGAUGAG X CGAA IAAGGCAA	3672
1969	CTTCTGAC T TCTTTCCT	1164	AGGAAAGA CUGAUGAG X CGAA IUCAGAAG	3673
1972	CTGACTTC T TTCCTTCT	1165	AGAAGGAA CUGAUGAG X CGAA IAAGUCAG	3674
1976	CTTCTTTC C TTCTATTC	1166	GAAUAGAA CUGAUGAG X CGAA IAAAGAAG	3675
1977	TTCTTTCC T TCTATTCG	1167	CGAAUAGA CUGAUGAG X CGAA IGAAAGAA	3676
1980	TTTCTTC T ATTGAGA	1168	UCUCGAAU CUGAUGAG X CGAA IAAGGAAA	3677
1991	TCGAGATC T CCTCGACA	1169	UGUCGAGG CUGAUGAG X CGAA IAUCUCGA	3678
1993	GAGATCTC C TCGACACC	1170	GGUGUCGA CUGAUGAG X CGAA IAGAUCUC	3679
1994	AGATCTCC T CGACACCG	1171	CGGUGUCG CUGAUGAG X CGAA IGAGAUCU	3680
1999	TCCTCGAC A CCGCTCT	1172	AGAGGCGG CUGAUGAG X CGAA IUCGAGGA	3681
2001	CTCGACAC C GCCTCTGC	1173	GCAGAGGC CUGAUGAG X CGAA IUGUCGAG	3682
2004	GACACCGC C TCTGCTCT	1174	AGAGCAGA CUGAUGAG X CGAA ICGGUGUC	3683
2005	ACACCGCC T CTGCTCTG	1175	CAGAGCAG CUGAUGAG X CGAA ICGGUGU	3684
2007	ACCGCCTC T GCTCTGTA	1176	UACAGAGC CUGAUGAG X CGAA IAGGCGGU	3685
2010	GCCTCTGC T CTGTATCG	1177	CGAUACAG CUGAUGAG X CGAA ICAGAGGC	3686
2012	CTCTGCTC T GTATCGGG	1178	CCCGAUAC CUGAUGAG X CGAA IAGCAGAG	3687
2025	CGGGGGGC C TTAGAGTC	1179	GACUCUAA CUGAUGAG X CGAA ICCCCCG	3688
2026	GGGGGGCC T TAGAGTCT	1180	AGACUCUA CUGAUGAG X CGAA IGCCCCC	3689
2034	TTAGAGTC T CCGGAACA	1181	UGUCCGG CUGAUGAG X CGAA IACUCUAA	3690
2036	AGAGTCTC C GGAACATT	1182	AAUGUCCG CUGAUGAG X CGAA IAGACUCU	3691
2042	TCCGGAAC A TTGTCAC	1183	GUGAACAA CUGAUGAG X CGAA IUUCCGGA	3692
2049	CATTGTTT C C CTCACCA	1184	UGGUGAGG CUGAUGAG X CGAA IAACAAUG	3693
2051	TTGTTTAC C TCACCATA	1185	UAUGGUGA CUGAUGAG X CGAA IUGAACAA	3694
2052	TGTTTACC T CACCATAC	1186	GUAUGGUG CUGAUGAG X CGAA IGUGAACAA	3695
2054	TTACCTC A CCATACCG	1187	CCGUAUGG CUGAUGAG X CGAA IAGGUGAA	3696
2056	CACCTCAC C ATACGGCA	1188	UGCCGUAU CUGAUGAG X CGAA IUGAGGUG	3697
2057	ACCTCACC A TACGGCAC	1189	GUGCCGUA CUGAUGAG X CGAA IGUGAGGU	3698
2064	CATACGGC A CTCAGGCA	1190	UGCCUGAG CUGAUGAG X CGAA ICCGUAUG	3699
2066	TACGGCAC T CAGGCAAG	1191	CUUGCCUG CUGAUGAG X CGAA IUGCCGUA	3700
2068	CGGCACTC A GGCAAGCT	1192	AGCUUGCC CUGAUGAG X CGAA IAGUGCCG	3701
2072	ACTCAGGC A AGCTATTC	1193	GAAUAGCU CUGAUGAG X CGAA ICCUGAGU	3702
2076	AGGCAAGC T ATTCTGTG	1194	CACAGAAU CUGAUGAG X CGAA ICUUGCCU	3703
2081	AGCTATTC T GTGTTGGG	1195	CCCAACAC CUGAUGAG X CGAA IAAUAGCU	3704
2105	GATGAATC T AGCCACCT	1196	AGGUGGCU CUGAUGAG X CGAA IAUUCAUC	3705
2109	AATCTAGC C ACCTGGGT	1197	ACCCAGGU CUGAUGAG X CGAA ICUAGAUA	3706
2110	ATCTAGCC A CCTGGGTG	1198	CACCCAGG CUGAUGAG X CGAA IGCUGAGU	3707
2112	CTAGCCAC C TGGGTGGG	1199	CCCACCCA CUGAUGAG X CGAA IUGGCUAG	3708
2113	TAGCCACC T GGGTGGGA	1200	UCCACCCC CUGAUGAG X CGAA IGUGGCUA	3709
2138	GGAAGATC C AGCATCCA	1201	UGGAUGCU CUGAUGAG X CGAA IAUCUUC	3710
2139	GAAGATCC A GCATCCAG	1202	CUGGAUGC CUGAUGAG X CGAA IGAUCUUC	3711
2142	GATCCAGC A TCCAGGGA	1203	UCCUGGA CUGAUGAG X CGAA ICUGGAUC	3712
2145	CCAGCATC C AGGGAATT	1204	AAUUCUU CUGAUGAG X CGAA IAUGCUGG	3713
2146	CAGCATCC A GGAATTA	1205	UAAUUCU CUGAUGAG X CGAA IGAUGCUG	3714
2161	TAGTAGTC A GCTATGTC	1206	GACAUAGC CUGAUGAG X CGAA IACUACUA	3715
2164	TAGTCAGC T ATGTCAAC	1207	GUUGACAU CUGAUGAG X CGAA ICUGACUA	3716

Table 38

2170	GCTATGTC A ACGTTAAT	1208	AUUAACGU CUGAUGAG X CGAA IACAUAGC	3717
2185	ATATGGGC C TAAAAATC	1209	GAUUUUUA CUGAUGAG X CGAA ICCCAUUAU	3718
2186	TATGGGCC T AAAAATCA	1210	UGAUUUUU CUGAUGAG X CGAA IGCCCAUA	3719
2194	TAAAAATC A GACAACTA	1211	UAGUUGUC CUGAUGAG X CGAA IAUUUUUA	3720
2198	AATCAGAC A ACTATTGT	1212	ACAAUAGU CUGAUGAG X CGAA IUCUGAUU	3721
2201	CAGACAAC T ATTGTGGT	1213	ACCACAAU CUGAUGAG X CGAA IUUGUCUG	3722
2213	GTGGTTTC A CATTTCTT	1214	AGGAAAUG CUGAUGAG X CGAA IAAACCAC	3723
2215	GGTTTCAC A TTCTCTGT	1215	ACAGGAAA CUGAUGAG X CGAA IUGAAACC	3724
2220	CACATTTT C TGTCTTAC	1216	GUAAGACA CUGAUGAG X CGAA IAAAUGUG	3725
2221	ACATTTTC T GTCTTACT	1217	AGUAAGAC CUGAUGAG X CGAA IGAAAUGU	3726
2225	TTCTGTGC T TACTTTTG	1218	CAAAAGUA CUGAUGAG X CGAA IACAGGAA	3727
2229	TGTCTTAC T TTTGGGCG	1219	CGCCCAAA CUGAUGAG X CGAA IUAAGACA	3728
2244	CGAGAAAC T GTTCTTGA	1220	UCAAGAAC CUGAUGAG X CGAA IUUUCUCG	3729
2249	AACTGTTC T TGAATATT	1221	AAUAUUCA CUGAUGAG X CGAA IAACAGUU	3730
2265	TTGGTGTG T TTTGGAGT	1222	ACUCCAAA CUGAUGAG X CGAA IACACCAA	3731
2284	GGATTGCG A CTCCTCCT	1223	AGGAGGAG CUGAUGAG X CGAA ICGAAUCC	3732
2286	ATTGCGAC T CCTCCTGC	1224	GCAGGAGG CUGAUGAG X CGAA IUGCGAAU	3733
2288	TCGCACTC C TCCTGCAT	1225	AUGCAGGA CUGAUGAG X CGAA IAGUGCGA	3734
2289	CGCACTCC T CCTGCATA	1226	UAUGCAGG CUGAUGAG X CGAA IGAGUGCG	3735
2291	CACTCCTC C TGCATATA	1227	UAUAUGCA CUGAUGAG X CGAA IAGGAGUG	3736
2292	ACTCCTCC T GCATATAG	1228	CUAUAUGC CUGAUGAG X CGAA IGAGGAGU	3737
2295	CCTCTGTC A TATAGACC	1229	GGUCUAUA CUGAUGAG X CGAA ICAGGAGG	3738
2303	ATATAGAC C ACCAAATG	1230	CAUUUGGU CUGAUGAG X CGAA IUCUAUUAU	3739
2304	TATAGACC A CCAAATGC	1231	GCAUUUGG CUGAUGAG X CGAA IGUCUAUA	3740
2306	TAGACCAC C AAATGCCC	1232	GGGCAUUU CUGAUGAG X CGAA IUGGUCUA	3741
2307	AGACCACC A AATGCCCC	1233	GGGGCAUU CUGAUGAG X CGAA IGUGGUCU	3742
2313	CCAAATGC C CCTATCTT	1234	AAGAUAGG CUGAUGAG X CGAA ICAUUUGG	3743
2314	CAAATGCC C CTATCTTA	1235	UAAGAUAG CUGAUGAG X CGAA IGCAUUUG	3744
2315	AAATGCCC C TATCTTAT	1236	AUAAGAUU CUGAUGAG X CGAA IGGCAUUU	3745
2316	AATGCCCC T ATCTTATC	1237	GAUAAGAU CUGAUGAG X CGAA IGGCAUUU	3746
2320	CCCCTATC T TATCAACA	1238	UGUUGAUA CUGAUGAG X CGAA IAUAGGGG	3747
2325	ATCTTATC A ACACTTCC	1239	GGAAGUGU CUGAUGAG X CGAA IAUAGGAU	3748
2328	TTATCAAC A CTCCGGA	1240	UCCGGAAG CUGAUGAG X CGAA IUUGAUAA	3749
2330	ATCAACAC T TCCGAAA	1241	UUUCCGGA CUGAUGAG X CGAA IUGUUGAU	3750
2333	AACACTTC C GGAAACTA	1242	UAGUUUCC CUGAUGAG X CGAA IAAGUGUU	3751
2340	CCGGAAAC T ACTGTTGT	1243	ACAACAGU CUGAUGAG X CGAA IUUUCCGG	3752
2343	GAAACTAC T GTTGTTAG	1244	CUAACAAC CUGAUGAG X CGAA IUAGUUUC	3753
2362	GAAGAGGC A GGTCCCCT	1245	AGGGGACC CUGAUGAG X CGAA ICCUCUUC	3754
2367	GGCAGGTC C CCTAGAAG	1246	CUUCUAGG CUGAUGAG X CGAA IACCUGCC	3755
2368	GCAGGTCC C CTAGAAGA	1247	UCUUCUAG CUGAUGAG X CGAA IGACCUGC	3756
2369	CAGGTCCC C TAGAAGAA	1248	UUCUUCUA CUGAUGAG X CGAA IGGACCUG	3757
2370	AGGTCCCC T AGAAGAAG	1249	CUUCUUCU CUGAUGAG X CGAA IGGACCUC	3758
2382	AGAAGAAC T CCCTCGCC	1250	GGCGAGGG CUGAUGAG X CGAA IUUCUUCU	3759
2384	AAGAACTC C CTCGCCTC	1251	GAGGCGAG CUGAUGAG X CGAA IAGUUCUU	3760
2385	AGAACTCC C TCGCCTCG	1252	CGAGGCGA CUGAUGAG X CGAA IGAGUUCU	3761
2386	GAACCTCC T CGCCTCGC	1253	GCGAGGCG CUGAUGAG X CGAA IGGAGUUC	3762
2390	TCCCTCGC C TCGCAGAC	1254	GUCUGCGA CUGAUGAG X CGAA ICGAGGGA	3763
2391	CCCTCGCC T CGCAGACG	1255	CGUCUGCG CUGAUGAG X CGAA IGCGAGGG	3764
2395	CGCCTCGC A GACGAAGG	1256	CCUUCGUC CUGAUGAG X CGAA ICGAGGCG	3765
2406	CGAAGGTC T CAATCGCC	1257	GGCGAUUG CUGAUGAG X CGAA IACCUUCG	3766
2408	AAGGTCTC A ATCGCCGC	1258	GCGGCGAU CUGAUGAG X CGAA IAGACCUU	3767

Table 38

2414	TCAATCGC C GCGTCGCA	1259	UGCGACGC CUGAUGAG X CGAA ICGAUUGA	3768
2422	CGCGTCGC A GAAGATCT	1260	AGAUCUUC CUGAUGAG X CGAA ICGACGCG	3769
2430	AGAAGATC T CAATCTCG	1261	CGAGAUUG CUGAUGAG X CGAA IAUUCUUCU	3770
2432	AAGATCTC A ATCTCGGG	1262	CCCAGAU CUGAUGAG X CGAA IAGAUCUU	3771
2436	TCTCAATC T CGGGAATC	1263	GAUUCCTG CUGAUGAG X CGAA IAUUGAGA	3772
2445	CGGGAATC T CAATGTTA	1264	UAACAUUG CUGAUGAG X CGAA IAUUCCCG	3773
2447	GGAATCTC A ATGTTAGT	1265	ACUAAAU CUGAUGAG X CGAA IAGAUUCC	3774
2460	TAGTATTC C TTGGACAC	1266	GUGUCCAA CUGAUGAG X CGAA IAAUACUA	3775
2461	AGTATTCC T TGGACACA	1267	UGUGUCCA CUGAUGAG X CGAA IGAAUACU	3776
2467	CCTTGGAC A CATAAGGT	1268	ACCUUUG CUGAUGAG X CGAA IUCCAAGG	3777
2469	TTGGACAC A TAAGGTGG	1269	CCACCUUA CUGAUGAG X CGAA IUGUCCAA	3778
2483	TGGGAAAC T TTACGGGG	1270	CCCCGUA CUGAUGAG X CGAA IUUCCCCA	3779
2493	TACGGGGC T TTATTCTT	1271	AAGAAUAA CUGAUGAG X CGAA ICCCCGUA	3780
2500	CTTTATTC T TCTACGGT	1272	ACCGUAGA CUGAUGAG X CGAA IAAUAAAG	3781
2503	TATTCTTC T ACGGTACC	1273	GGUACCGU CUGAUGAG X CGAA IAAGAAUA	3782
2511	TACGGTAC C TTGCTTTA	1274	UAAAGCAA CUGAUGAG X CGAA IUACCGUA	3783
2512	ACGGTACC T TGCTTTAA	1275	UUAAGCA CUGAUGAG X CGAA IGUACCGU	3784
2516	TACCTTGC T TTAATCCT	1276	AGGAUUAA CUGAUGAG X CGAA ICAAGGUA	3785
2523	CTTTAATC C TAAATGGC	1277	GCCAUUUA CUGAUGAG X CGAA IAUUAAAG	3786
2524	TTTAATCC T AAATGGCA	1278	UGCCAUUU CUGAUGAG X CGAA IGAUUAUA	3787
2532	TAAATGGC A AACTCCTT	1279	AAGGAGUU CUGAUGAG X CGAA ICCAUUUA	3788
2536	TGGCAAAC T CCTTCTTT	1280	AAAGAAGG CUGAUGAG X CGAA IUUUGCCA	3789
2538	GCAAACCT C TTCTTTTC	1281	GAAAAGAA CUGAUGAG X CGAA IAGUUUGC	3790
2539	CAAACCTC T TCTTTTCC	1282	GGAAAAGA CUGAUGAG X CGAA IGAGUUUG	3791
2542	ACTCCTTC T TTTCTTGA	1283	UCAGGAAA CUGAUGAG X CGAA IAAGGAGU	3792
2547	TTCTTTTC C TGACATTC	1284	GAAUGUCA CUGAUGAG X CGAA IAAAAGAA	3793
2548	TCTTTTCC T GACATTCA	1285	UGAUGUC CUGAUGAG X CGAA IGAAAAGA	3794
2552	TTCTTGAC A TTCATTG	1286	CAAUGAA CUGAUGAG X CGAA IUCAGGAA	3795
2556	TGACATTC A TTTGCAGG	1287	CCUGCAA CUGAUGAG X CGAA IAAUGUCA	3796
2562	TCATTGAC A GGAGGACA	1288	UGUCCUCC CUGAUGAG X CGAA ICAAUGA	3797
2570	AGGAGGAC A TTGTTGAT	1289	AUCAACA CUGAUGAG X CGAA IUCCUCCU	3798
2589	ATGTAAGC A ATTTGTGG	1290	CCACAAU CUGAUGAG X CGAA ICUUACAU	3799
2601	TGTGGGGC C CTTACAG	1291	CUGUAAG CUGAUGAG X CGAA ICCCCACA	3800
2602	GTGGGGCC C CTTACAGT	1292	ACUGUAAG CUGAUGAG X CGAA IGCCCCAC	3801
2603	TGGGGCCC C TTACAGTA	1293	UACUGUA CUGAUGAG X CGAA IGGCCCCA	3802
2604	GGGGCCCC T TACAGTAA	1294	UUACUGUA CUGAUGAG X CGAA IGGGCCCC	3803
2608	CCCCTTAC A GTAAATGA	1295	UCAUUUAC CUGAUGAG X CGAA IUAAGGGG	3804
2621	ATGAAAAC A GGAGACTT	1296	AAGUCUCC CUGAUGAG X CGAA IUUUUCAU	3805
2628	CAGGAGAC T TAAATTAA	1297	UUAUUUA CUGAUGAG X CGAA IUCUCCUG	3806
2638	AAATTAAC T ATGCCTGC	1298	GCAGGCAU CUGAUGAG X CGAA IUUAAUUU	3807
2643	AACTATGC C TGCTAGGT	1299	ACCUAGCA CUGAUGAG X CGAA ICAUAGUU	3808
2644	ACTATGCC T GCTAGGTT	1300	AACCUAGC CUGAUGAG X CGAA IGCAUAGU	3809
2647	ATGCCTGC T AGGTTTTA	1301	UAAACCU CUGAUGAG X CGAA ICAGGCAU	3810
2658	GTTTTATC C CAATGTTA	1302	UAACAUUG CUGAUGAG X CGAA IAUAAAAC	3811
2659	TTTTATCC C AATGTTAC	1303	GUAACAU CUGAUGAG X CGAA IGAUAAAA	3812
2660	TTTATCCC A ATGTTACT	1304	AGUAACAU CUGAUGAG X CGAA IGGUAAAA	3813
2668	AATGTTAC T AAATATTT	1305	AAAUUUU CUGAUGAG X CGAA IUAAAUUU	3814
2679	ATATTTGC C CTTAGATA	1306	UAUCUAAG CUGAUGAG X CGAA ICAAUUUU	3815
2680	TATTTGCC C TTAGATAA	1307	UUUUCUA CUGAUGAG X CGAA IGCAAUUA	3816
2681	ATTTGCC C TTAGATAA	1308	UUUUCUA CUGAUGAG X CGAA IGGCAUUU	3817
2696	AAGGGATC A AACCGTAT	1309	AUACGGUU CUGAUGAG X CGAA IAUCCCUU	3818

Table 38

2700	GATCAAAC C GTATTATC	1310	GAUAAUAC CUGAUGAG X CGAA IUUUGAUC	3819
2709	GTATTATC C AGAGTATG	1311	CAUACUCU CUGAUGAG X CGAA IAUAAUAC	3820
2710	TATTATCC A GAGTATGT	1312	ACAUACUC CUGAUGAG X CGAA IGAUAAUA	3821
2727	AGTTAATC A TTAATTCC	1313	GGAAGUAA CUGAUGAG X CGAA IAUUAACU	3822
2732	ATCATTAC T TCCAGACG	1314	CGUCUGGA CUGAUGAG X CGAA IUAAUGAU	3823
2735	ATTACTTC C AGACGCGA	1315	UCGCGUCU CUGAUGAG X CGAA IAAGUAAU	3824
2736	TTACTTCC A GACGCGAC	1316	GUCGCGUC CUGAUGAG X CGAA IGAAGUAA	3825
2745	GACGCGAC A TTATTTAC	1317	GUAAAUAA CUGAUGAG X CGAA IUCGCGUC	3826
2754	TTATTTAC A CACTCTTT	1318	AAAGAGUG CUGAUGAG X CGAA IUAAAUAA	3827
2756	ATTTACAC A CTCTTGG	1319	CCAAAGAG CUGAUGAG X CGAA IUGUAAAU	3828
2758	TTACACAC T CTTTGGAA	1320	UUCCAAAG CUGAUGAG X CGAA IUGUGUAA	3829
2760	ACACACTC T TTGGAAGG	1321	CCUCCAA CUGAUGAG X CGAA IAGUGUGU	3830
2777	CGGGGATC T TATATAAA	1322	UUUAUUA CUGAUGAG X CGAA IAUCCCCG	3831
2794	AGAGAGTC C ACACGTAG	1323	CUACGUGU CUGAUGAG X CGAA IACUCUCU	3832
2795	GAGAGTCC A CACGTAGC	1324	GCUACGUG CUGAUGAG X CGAA IGACUCUC	3833
2797	GAGTCCAC A CGTAGCGC	1325	GCGCUACG CUGAUGAG X CGAA IUGGACUC	3834
2806	CGTAGCGC C TCATTTTG	1326	CAAAUGA CUGAUGAG X CGAA ICGCUACG	3835
2807	GTAGCGCC T CATTTTGC	1327	GCAAAUG CUGAUGAG X CGAA ICGCGUAC	3836
2809	AGCGCCTC A TTTTGCGG	1328	CCGCAAAA CUGAUGAG X CGAA IAGGCGCU	3837
2821	TGCGGGTC A CCATATTC	1329	GAAUAUGG CUGAUGAG X CGAA IACCCGCA	3838
2823	CGGGTCAC C ATATTCTT	1330	AAGAAUUA CUGAUGAG X CGAA IUGACCCG	3839
2824	GGGTCACC A TATTCTTG	1331	CAAGAAUA CUGAUGAG X CGAA IGUGACCC	3840
2830	CCATATTC T TGGGAACA	1332	UGUCCCA CUGAUGAG X CGAA IAAUAUGG	3841
2838	TTGGGAAC A AGATCTAC	1333	GUAGAUCU CUGAUGAG X CGAA IUUCCCAA	3842
2844	ACAAGATC T ACAGCATG	1334	CAUGCUGU CUGAUGAG X CGAA IAUUCUGU	3843
2847	AGATCTAC A GCATGGGA	1335	UCCCAUG CUGAUGAG X CGAA IUAGAUCU	3844
2850	TCTACAGC A TGGGAGGT	1336	ACCUCCA CUGAUGAG X CGAA ICUGUAGA	3845
2864	GGTTGGTC T TCCAAACC	1337	GGUUUGGA CUGAUGAG X CGAA IACCAACC	3846
2867	TGGTCTTC C AAACCTCG	1338	CGAGGUUU CUGAUGAG X CGAA IAAGACCA	3847
2868	GGTCTTCC A AACCTCGA	1339	UCGAGGUU CUGAUGAG X CGAA IGAAGACC	3848
2872	TTCCAAAC C TCGAAAAG	1340	CUUUUCGA CUGAUGAG X CGAA IUUUGGAA	3849
2873	TCCAAACC T CGAAAAGG	1341	CCUUUUCG CUGAUGAG X CGAA IGUUUGGA	3850
2883	GAAAAGGC A TGGGGACA	1342	UGUCCCA CUGAUGAG X CGAA ICCUUUUC	3851
2891	ATGGGGAC A AATCTTTC	1343	GAAAGAUU CUGAUGAG X CGAA IUCCCCAU	3852
2896	GACAAATC T TTCTGTCC	1344	GGACAGAA CUGAUGAG X CGAA IAUUUGUC	3853
2900	AATCTTTC T GTCCCCAA	1345	UUGGGGAC CUGAUGAG X CGAA IAAAGAUU	3854
2904	TTTCTGTC C CCAATCCC	1346	GGGAUUGG CUGAUGAG X CGAA IACAGAAA	3855
2905	TTCTGTCC C CAATCCCC	1347	GGGAUUG CUGAUGAG X CGAA IGACAGAA	3856
2906	TCTGTCCC C AATCCCTT	1348	AGGGGAUU CUGAUGAG X CGAA IGGACAGA	3857
2907	CTGTCCCC A ATCCCCTG	1349	CAGGGGAU CUGAUGAG X CGAA IGGACAG	3858
2911	CCCCAATC C CTGGGAT	1350	AUCCAGG CUGAUGAG X CGAA IAUUGGGG	3859
2912	CCCAATCC C CTGGGATT	1351	AAUCCAG CUGAUGAG X CGAA IGAUUGGG	3860
2913	CCAATCCC C TGGGATTCT	1352	GAAUCCCA CUGAUGAG X CGAA IGGAUUGG	3861
2914	CAATCCCC T GGGATTCT	1353	AGAAUCCC CUGAUGAG X CGAA IGGAUUG	3862
2922	TGGGATTCT T TCCCCGAT	1354	AUCGGGGA CUGAUGAG X CGAA IAAUCCCA	3863
2925	GATTCTTC C CGATCAT	1355	AUGAUCGG CUGAUGAG X CGAA IAAGAAUC	3864
2926	ATTCTTCC C CGATCATC	1356	GAUGAUCG CUGAUGAG X CGAA IGAAGAAU	3865
2927	TTCTTCCC C GATCATCA	1357	UGAUGAUC CUGAUGAG X CGAA IGGAGAA	3866
2932	CCCCGATC A TCAGTTGG	1358	CCAACUGA CUGAUGAG X CGAA IAUCCGGG	3867
2935	CGATCATC A GTTGGACC	1359	GGUCCAAC CUGAUGAG X CGAA IAUGAUCG	3868
2943	AGTTGGAC C CTGCATTCT	1360	GAAUGCAG CUGAUGAG X CGAA IUCCAACU	3869

Table 38

2944	GTTGGACC C TGCATTCA	1361	UGAAUGCA CUGAUGAG X CGAA IGUCCAAC	3870
2945	TTGGACCC T GCATTCAA	1362	UUGAAUGC CUGAUGAG X CGAA IGGUCCAA	3871
2948	GACCTGCA A TTCAAAGC	1363	GCUUUGAA CUGAUGAG X CGAA ICAGGGUC	3872
2952	CTGCATT C AAGCCAAC	1364	GUUGGCUU CUGAUGAG X CGAA IAAUGCAG	3873
2957	TTCAAAGC C AACTCAGT	1365	ACUGAGUU CUGAUGAG X CGAA ICUUUGAA	3874
2958	TCAAAGCC A ACTCAGTA	1366	UACUGAGU CUGAUGAG X CGAA IGCUUUGA	3875
2961	AAGCCAAC T CAGTAAAT	1367	AUUUACUG CUGAUGAG X CGAA IUUGGCUU	3876
2963	GCCAATC A GTAAATCC	1368	GGAUUUAC CUGAUGAG X CGAA IAGUUGGC	3877
2971	AGTAAATC C AGATTGGG	1369	CCCAAUCU CUGAUGAG X CGAA IAUUUACU	3878
2972	GTAAATCC A GATTGGGA	1370	UCCCAAUC CUGAUGAG X CGAA IGAUUUAC	3879
2982	ATTGGGAC C TCAACCCG	1371	CGGGUUGA CUGAUGAG X CGAA IUCCCAAU	3880
2983	TTGGGACC T CAACCCGC	1372	GCGGGUUG CUGAUGAG X CGAA IGUCCCAA	3881
2985	GGGACCTC A ACCCGCAC	1373	GUGCGGUU CUGAUGAG X CGAA IAGGUCCC	3882
2988	ACCTCAAC C CGCACAAG	1374	CUUGUGCG CUGAUGAG X CGAA IUUGAGGU	3883
2989	CCTCAACC C GCACAAGG	1375	CCUUGUGC CUGAUGAG X CGAA IGUUGAGG	3884
2992	CAACCCGC A CAAGGACA	1376	UGUCCUUG CUGAUGAG X CGAA ICGGGUUG	3885
2994	ACCCGCAC A AGGACAAC	1377	GUUGUCCU CUGAUGAG X CGAA IUGCGGUU	3886
3000	ACAAGGAC A ACTGGCCG	1378	CGGCCAGU CUGAUGAG X CGAA IUCCUUGU	3887
3003	AGGACAAC T GGCCGGAC	1379	GUCCGGCC CUGAUGAG X CGAA IUUGUCCU	3888
3007	CAACTGGC C GGACGCCA	1380	UGGCGUCC CUGAUGAG X CGAA ICCAGUUG	3889
3014	CCGACGCG C AACAAGGT	1381	ACCUUGUU CUGAUGAG X CGAA ICGUCCGG	3890
3015	CGGACGCC A ACAAGGTG	1382	CACCUUGU CUGAUGAG X CGAA ICGUCCGG	3891
3018	ACGCCAAC A AGGTGGGA	1383	UCCCACCU CUGAUGAG X CGAA IUUGGCGU	3892
3035	GTGGGAGC A TTCGGGCC	1384	GGCCCGAA CUGAUGAG X CGAA ICUCCAC	3893
3043	ATTCGGGC C AGGGTTCA	1385	UGAACCCU CUGAUGAG X CGAA ICCCGAAU	3894
3044	TTCGGGCC A GGGTTTCA	1386	GUGAACCC CUGAUGAG X CGAA IGCCCGAA	3895
3051	CAGGGTTC A CCCCTCCC	1387	GGGAGGGG CUGAUGAG X CGAA IAACCCUG	3896
3053	GGGTTCAC C CCTCCCCA	1388	UGGGGAGG CUGAUGAG X CGAA IUGAACCC	3897
3054	GGTTCACC C CTCCCAT	1389	AUGGGGAG CUGAUGAG X CGAA IGUGAAC	3898
3055	GTTCACCC C TCCCATG	1390	CAUGGGGA CUGAUGAG X CGAA IGGUGAAC	3899
3056	TTCACCCC C CCCCATGG	1391	CCAUGGGG CUGAUGAG X CGAA IGGUGAA	3900
3058	CACCCCTC C CCATGGGG	1392	CCCCAUGG CUGAUGAG X CGAA IAGGGGUG	3901
3059	ACCCCTCC C CATGGGGG	1393	CCCCCAUG CUGAUGAG X CGAA IAGGGGUG	3902
3060	CCCCTCCC C ATGGGGGA	1394	UCCCCCAU CUGAUGAG X CGAA IGGAGGGG	3903
3061	CCCTCCCC A TGGGGGAC	1395	GUCCCCCA CUGAUGAG X CGAA IGGAGGGG	3904
3070	TGGGGGAC T GTTGGGGT	1396	ACCCCAAC CUGAUGAG X CGAA IUCCCCCA	3905
3084	GGTGGAGC C CTCACGCT	1397	AGCGUGAG CUGAUGAG X CGAA ICUCCACC	3906
3085	GTGGAGCC C TCACGCTC	1398	GAGCGUGA CUGAUGAG X CGAA IGCUCCAC	3907
3086	TGGAGCCC T CACGCTCA	1399	UGAGCGUG CUGAUGAG X CGAA IGGCUCCA	3908
3088	GAGCCCTC A CGCTCAGG	1400	CCUGAGCG CUGAUGAG X CGAA IAGGGCUC	3909
3092	CCTCACGC T CAGGGCCT	1401	AGGCCCUG CUGAUGAG X CGAA ICGUGAGG	3910
3094	TCACGCTC A GGGCCTAC	1402	GUAGGCCC CUGAUGAG X CGAA IAGCGUGA	3911
3099	CTCAGGGC C TACTCACA	1403	UGUGAGUA CUGAUGAG X CGAA ICCUGAG	3912
3100	TCAGGGCC T ACTCACA	1404	UUGUGAGU CUGAUGAG X CGAA IGCCUGA	3913
3103	GGGCCTAC T CACAACCTG	1405	CAGUUGUG CUGAUGAG X CGAA IUAGGCCC	3914
3105	GCCTACTC A CACTGTG	1406	CACAGUUG CUGAUGAG X CGAA IAGUAGGC	3915
3107	CTACTCAC A ACTGTGCC	1407	GGCACAGU CUGAUGAG X CGAA IUAGUAG	3916
3110	CTCACAAC T GTGCCAGC	1408	GCUUGGAC CUGAUGAG X CGAA IUUGUGAG	3917
3115	AACTGTGC C AGCAGCTC	1409	GAGCUGCU CUGAUGAG X CGAA ICACAGUU	3918
3116	ACTGTGCC A GCAGCTCC	1410	GGAGCUGC CUGAUGAG X CGAA IGCACAGU	3919
3119	GTGCCAGC A GCTCCTCC	1411	GGAGGAGC CUGAUGAG X CGAA ICUGGCAC	3920

Table 38

3122	CCAGCAGC T CCTCCTCC	1412	GGAGGAGG CUGAUGAG X CGAA ICUGCUGG	3921
3124	AGCAGCTC C TCCTCCTG	1413	CAGGAGGA CUGAUGAG X CGAA IAGCUGCU	3922
3125	GCAGCTCC T CCTCCTGC	1414	GCAGGAGG CUGAUGAG X CGAA IGAGCUGC	3923
3127	AGCTCCTC C TCCTGCCT	1415	AGGCAGGA CUGAUGAG X CGAA IAGGAGCU	3924
3128	GCTCCTCC T CCTGCCTC	1416	GAGGAGG CUGAUGAG X CGAA IGAGGAGC	3925
3130	TCCTCCTC C TGCCTCCA	1417	UGGAGGCA CUGAUGAG X CGAA IAGGAGGA	3926
3131	CCTCCTCC T GCCTCCAC	1418	GUGGAGGC CUGAUGAG X CGAA IGAGGAGG	3927
3134	CCTCCTGC C TCCACCAA	1419	UUGGUGGA CUGAUGAG X CGAA ICAGGAGG	3928
3135	CTCCTGCC T CCACCAAT	1420	AUUGGUGG CUGAUGAG X CGAA IGCAGGAG	3929
3137	CCTGCCTC C ACCAATCG	1421	CGAUUGGU CUGAUGAG X CGAA IAGGCAGG	3930
3138	CTGCCTCC A CCAATCGG	1422	CCGAUUGG CUGAUGAG X CGAA IGAGGCAG	3931
3140	GCCTCCAC C AATCGGCA	1423	UGCCGAUU CUGAUGAG X CGAA IUGGAGGC	3932
3141	CCTCCACC A ATCGGCAG	1424	CUGCCGAU CUGAUGAG X CGAA IGUGGAGG	3933
3148	CAATCGGC A GTCAGGAA	1425	UUCCUGAC CUGAUGAG X CGAA ICCGAUUG	3934
3152	CGGCAGTC A GGAAGGCA	1426	UGCCUUC CUGAUGAG X CGAA IACUGCCG	3935
3160	AGGAAGGC A GCCTACTC	1427	GAGUAGGC CUGAUGAG X CGAA ICCUUCU	3936
3163	AAGGCAGC C TACTCCCT	1428	AGGGAGUA CUGAUGAG X CGAA ICUGCCUU	3937
3164	AGGCAGCC T ACTCCCTT	1429	AAGGGAGU CUGAUGAG X CGAA IGCUGCCU	3938
3167	CAGCTAC T CCCTTATC	1430	GAUAAGGG CUGAUGAG X CGAA IUAGGCUG	3939
3169	GCCTACTC C CTTATCTC	1431	GAGAUAA CUGAUGAG X CGAA IAGUAGGC	3940
3170	CCTACTCC C TTATCTCC	1432	GGAGAUAA CUGAUGAG X CGAA IGAGUAGG	3941
3171	CTACTCCC T TATCTCCA	1433	UGGAGUA CUGAUGAG X CGAA IGGAGUAG	3942
3176	CCCTTATC T CCACCTCT	1434	AGAGGUGG CUGAUGAG X CGAA IAUAGGG	3943
3178	CTTATCTC C ACCTCTAA	1435	UUAGAGGU CUGAUGAG X CGAA IAGAUAG	3944
3179	TTATCTCC A CCTCTAAG	1436	CUUAGAGG CUGAUGAG X CGAA IGAGAUAA	3945
3181	ATCTCCAC C TCTAAGGG	1437	CCCUUAGA CUGAUGAG X CGAA IUUGAGAU	3946
3182	TCTCCACC T CTAAGGGA	1438	UCCCUAG CUGAUGAG X CGAA IGUGGAGA	3947
3184	TCCACCTC T AAGGGACA	1439	UGUCCUU CUGAUGAG X CGAA IAGGUGGA	3948
3192	TAAGGGAC A CTCATCCT	1440	AGGAUGAG CUGAUGAG X CGAA IUCCCUUA	3949
3194	AGGGACAC T CATCCTCA	1441	UGAGGAUG CUGAUGAG X CGAA IUUGCCCU	3950
3196	GGACACTC A TCCTCAGG	1442	CCUGAGGA CUGAUGAG X CGAA IAGUGUCC	3951
3199	CACTCATC C TCAGGCCA	1443	UGGCCUGA CUGAUGAG X CGAA IAUGAGUG	3952
3200	ACTCATCC T CAGGCCAT	1444	AUGGCCUG CUGAUGAG X CGAA IGAUGAGU	3953
3202	TCATCCTC A GGCCATGC	1445	GCAUGGCC CUGAUGAG X CGAA IAGGAUGA	3954
3206	CCTCAGGC C ATGCAGTG	1446	CACUGCAU CUGAUGAG X CGAA ICCUGAGG	3955
3207	CTCAGGCC A TGCAGTGG	1447	CCACUGCA CUGAUGAG X CGAA IGCCUGAG	3956

Input Sequence = AF100308. Cut Site = CH/.

Stem Length = 8 . Core Sequence = CUGAUGAG X CGAA (X = GCCGUUAGGC or other stem II)
AF100308 (Hepatitis B virus strain 2-18, 3215 bp)

Table 39

Table 39: Human HBV G-cleaver Ribozyme and Substrate Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
61	ACUUUCCU G CUGGUGGC	1448	GCCACCAG UGAUG GCAUGCACUAUGC GCG AGGAAAGU	3957
87	GGAACAGU G AGCCUUGC	1449	GCAGGCU UGAUG GCAUGCACUAUGC GCG ACUGUUC	3958
94	UGAGCCCU G CUCAGAAU	1450	AUUCUGAG UGAUG GCAUGCACUAUGC GCG AGGCUCA	3959
112	CUGUCUCU G CCAUAUCG	1451	CGAUAUGG UGAUG GCAUGCACUAUGC GCG AGAGACAG	3960
132	AUCUUAUC G AAGACUGG	1452	CCAGUCUU UGAUG GCAUGCACUAUGC GCG GAUAAGAU	3961
153	CCUGUACC G AACAUUGA	1453	UCCAUGUU UGAUG GCAUGCACUAUGC GCG GGUACAGG	3962
169	AGAACAU G CAUCAGGA	1454	UCCUGAUG UGAUG GCAUGCACUAUGC GCG GAUGUUCU	3963
192	GGACCCCU G CUCGUGUU	1455	AACACGAG UGAUG GCAUGCACUAUGC GCG AGGGUCC	3964
222	UUCUGUU G ACAAAGAA	1456	AUUUUUU UGAUG GCAUGCACUAUGC GCG AACAGAA	3965
315	CAAAAUUC G CAGUCCCA	1457	UGGGACUG UGAUG GCAUGCACUAUGC GCG GAUUUUUG	3966
374	UGGUUAUC G CUGGAUGU	1458	ACAUCAG UGAUG GCAUGCACUAUGC GCG GAUAACCA	3967
387	AUGUGUCU G CGGCGUUU	1459	AAACGCCG UGAUG GCAUGCACUAUGC GCG AGACACAU	3968
410	CUUCCUCU G CAUCCUUC	1460	GCAGGAUG UGAUG GCAUGCACUAUGC GCG AGAGGAAG	3969
417	UGCAUCCU G CUGCUAUG	1461	CAUAGCAG UGAUG GCAUGCACUAUGC GCG AGGAUGCA	3970
420	AUCCUGCU G CUAUGCCU	1462	AGGCAUAG UGAUG GCAUGCACUAUGC GCG AGCAGGAU	3971
425	GCUGCUAU G CCUCAUCU	1463	AGAUGAGG UGAUG GCAUGCACUAUGC GCG AUAGCAGC	3972
468	GGUAUGUU G CCGGUUUG	1464	CAACCGGG UGAUG GCAUGCACUAUGC GCG AACAUACC	3973
518	CGGACCAU G CAAAACCU	1465	AGGUUUUG UGAUG GCAUGCACUAUGC GCG AUGGUCCG	3974
527	CAAAACCU G CACAACUC	1466	GAGUUUGU UGAUG GCAUGCACUAUGC GCG AGGUUUUG	3975
538	CAACUCCU G CUCAAAGG	1467	UCCUUUGAG UGAUG GCAUGCACUAUGC GCG AGGAGUUG	3976
569	CUCAUGUU G CUGUACAA	1468	UUGUACAG UGAUG GCAUGCACUAUGC GCG AACAUAG	3977
596	CGGAACU G CACCUUGA	1469	UACAGGUG UGAUG GCAUGCACUAUGC GCG AGUUUCCG	3978
631	GGGCUUUC G CAAAUAAC	1470	GUUUUUUG UGAUG GCAUGCACUAUGC GCG GAAAGCCC	3979
687	UUACUAGU G CCAUUUGU	1471	ACAAAUUG UGAUG GCAUGCACUAUGC GCG ACUAGUAA	3980
747	AUAUGGAU G AUGUGGUU	1472	AACCAUUA UGAUG GCAUGCACUAUGC GCG AUCCAUAU	3981
783	AACAUCUU G AGUCCCUU	1473	AAGGACU UGAUG GCAUGCACUAUGC GCG AAGAUGUU	3982
795	CCCUUUUU G CCGCUGUU	1474	AACAGCGG UGAUG GCAUGCACUAUGC GCG AUAAAGGG	3983
798	UUUAUGCC G CUGUUACC	1475	GGUAACAG UGAUG GCAUGCACUAUGC GCG GGCAUAAA	3984
911	GGCACAUU G CCACAGGA	1476	UCCUGUGG UGAUG GCAUGCACUAUGC GCG AAUGUGCC	3985
978	GGCCUAUU G AUUGGAAA	1477	UUUCCAUA UGAUG GCAUGCACUAUGC GCG AAUAGGCC	3986

Table 39

997	AUGUCAAC G AAUUGUGG	1478	CCACAAU UGAUG GCAUGCACUAUGC GCG GUUGACAU	3987
1020	UGGGGUU G CCGCCCU	1479	AGGGGGG UGAUG GCAUGCACUAUGC GCG AAACCCCA	3988
1023	GGUUGCC G CCCUUC	1480	GAAAGGG UGAUG GCAUGCACUAUGC GCG GGCAAACC	3989
1034	CCUUCAC G CAAUGUGG	1481	CCACAUU UGAUG GCAUGCACUAUGC GCG GUGAAAGG	3990
1050	GAUAUUCU G CUUUAUUG	1482	CAUUAAG UGAUG GCAUGCACUAUGC GCG AGAAUAUC	3991
1058	GCUUAAU G CCUUAUA	1483	UAUAAAG UGAUG GCAUGCACUAUGC GCG AUUAAAGC	3992
1068	CUUUAU G CAUGCAUA	1484	UAUGCAU UGAUG GCAUGCACUAUGC GCG AUUAAAG	3993
1072	AUAUGCAU G CAUACAAG	1485	CUUGUUG UGAUG GCAUGCACUAUGC GCG AUGCAUUA	3994
1103	ACUUCUC G CCAACUUA	1486	UAAUUGG UGAUG GCAUGCACUAUGC GCG GAGAAAGU	3995
1139	CAGUAUGU G AACCUUUA	1487	UAAAGUU UGAUG GCAUGCACUAUGC GCG ACAUACUG	3996
1155	ACCCGGU G CUCGGCAA	1488	UUGCCGAG UGAUG GCAUGCACUAUGC GCG AACGGGUU	3997
1177	UGGUCUUAU G CCAAGUGU	1489	ACAUUGG UGAUG GCAUGCACUAUGC GCG AUAGACCA	3998
1188	AAGUGUU G CUGACGCA	1490	UGCUCAG UGAUG GCAUGCACUAUGC GCG AAACACUU	3999
1191	UGUUGCU G ACGCAACC	1491	GGUUGCU UGAUG GCAUGCACUAUGC GCG AGCAAACA	4000
1194	UUGUGAC G CAACCCCC	1492	GGGGUUG UGAUG GCAUGCACUAUGC GCG GUCAGCAA	4001
1234	CCAUCAGC G CAUGCGUG	1493	CACGAUG UGAUG GCAUGCACUAUGC GCG GCUGAUGG	4002
1238	CAGCGCAU G CGUGGAAC	1494	GUUCCAG UGAUG GCAUGCACUAUGC GCG AUGCGCUG	4003
1262	UCUCCUCU G CCGAUCCA	1495	UGGAUCG UGAUG GCAUGCACUAUGC GCG AGAGGAGA	4004
1265	CCUCUGCC G AUCCAUAU	1496	GUUAGAU UGAUG GCAUGCACUAUGC GCG GGCAGAGG	4005
1275	UCCAUAAC G CGGAACUC	1497	GAGUCCG UGAUG GCAUGCACUAUGC GCG GGUUAGGA	4006
1290	UCCUAGCC G CUUGUUUU	1498	AAAACAAG UGAUG GCAUGCACUAUGC GCG GGUUAGGA	4007
1299	CUUGUUU G CUCGCAGC	1499	GCUGCGAG UGAUG GCAUGCACUAUGC GCG AAAACAAG	4008
1303	UUUUGUC G CAGCAGGU	1500	ACUUGCU UGAUG GCAUGCACUAUGC GCG GAGCAAAA	4009
1335	UCGGGACU G ACAAUUCU	1501	AGAAUUGU UGAUG GCAUGCACUAUGC GCG AGUCCCGA	4010
1349	UCUGUCGU G CUCUCCCG	1502	CGGGAGAG UGAUG GCAUGCACUAUGC GCG ACGACAGA	4011
1357	GCUCUCCC G CAAUAUA	1503	UAUAUUG UGAUG GCAUGCACUAUGC GCG GGGAGAGC	4012
1382	CCAUGGCU G CUAGGCUG	1504	CAGCUAG UGAUG GCAUGCACUAUGC GCG AGCCAUGG	4013
1392	UAGGUCUGU G CUGCCAAC	1505	GUUGGCAG UGAUG GCAUGCACUAUGC GCG ACAGCCUA	4014
1395	GCUGUGCU G CCAACUGG	1506	CCAGUUGG UGAUG GCAUGCACUAUGC GCG AGCACAGC	4015
1411	GAUCUAC G CGGGACGU	1507	ACGUCCCG UGAUG GCAUGCACUAUGC GCG GUAGGAUC	4016
1442	CCGUCGGC G CUGAAUCC	1508	GGAUUCAG UGAUG GCAUGCACUAUGC GCG GCCGACGG	4017
1445	UCGGCGCU G AAUCCCGC	1509	GCGGGAU UGAUG GCAUGCACUAUGC GCG AGCGCCGA	4018
1452	UGAAUCCC G CGGACGAC	1510	GUCGUCCG UGAUG GCAUGCACUAUGC GCG GGGAUJCA	4019
1458	CCGCGGAC G ACCCCUCC	1511	GGAGGGGU UGAUG GCAUGCACUAUGC GCG GUCCGCGG	4020

Table 39

1474	CCGGGGCC G CUUGGGC	1512	GCCCAAG UGAUG GCAUGCACUAUGC GCG GCGCCCGG	4021
1489	GCUCUACC G CCCGCUUC	1513	GAAGCGG UGAUG GCAUGCACUAUGC GCG GGUAGAGC	4022
1493	UACCGCC G CUUCUCG	1514	CGGAAAG UGAUG GCAUGCACUAUGC GCG GCGCGGUA	4023
1501	GCUCUCC G CCUAUUG	1515	ACAAAGG UGAUG GCAUGCACUAUGC GCG GGAGAAGC	4024
1513	AUUGUACC G ACCGUCCA	1516	UGGACGGU UGAUG GCAUGCACUAUGC GCG GGUACAAU	4025
1528	CACGGGC G CACCUCUC	1517	GAGAGGUG UGAUG GCAUGCACUAUGC GCG GCCCCGUG	4026
1542	CUCUUAC G CGGACUC	1518	GGAGUCG UGAUG GCAUGCACUAUGC GCG GUAAAGAG	4027
1559	CCGUCUG G CCUUCUA	1519	UGAGAAGG UGAUG GCAUGCACUAUGC GCG ACAGACGG	4028
1571	UCUCAUCU G CCGGACCG	1520	CGGUCGG UGAUG GCAUGCACUAUGC GCG AGAUGAGA	4029
1583	GACCGUG G CACUUCGC	1521	GCGAAGUG UGAUG GCAUGCACUAUGC GCG ACACGGUC	4030
1590	UGCACUUC G CUUCACCU	1522	AGGUGAAG UGAUG GCAUGCACUAUGC GCG GAAUGGCA	4031
1601	UCACCUUC G CACGUCGC	1523	GCGACGUG UGAUG GCAUGCACUAUGC GCG AGAGGUGA	4032
1608	UGCACGUC G CAUGGAGA	1524	UCUCCAUG UGAUG GCAUGCACUAUGC GCG GACGUGCA	4033
1624	ACCACCGU G AAGGCCA	1525	UGGGCGUU UGAUG GCAUGCACUAUGC GCG ACGGUGGU	4034
1628	CCGUGAAC G CCCACAGG	1526	CCUGUGGG UGAUG GCAUGCACUAUGC GCG GUUCACGG	4035
1642	AGGAACCU G CCCAAGGU	1527	ACCUUGGG UGAUG GCAUGCACUAUGC GCG AGGUUCCU	4036
1654	AAGGUUU G CAUAAGAG	1528	CUCUUUUG UGAUG GCAUGCACUAUGC GCG AAGACCUU	4037
1690	AUGUCAAC G ACCGACCU	1529	AGGUCGGU UGAUG GCAUGCACUAUGC GCG GUUGACAU	4038
1694	CAACGACC G ACCUUGAG	1530	CUCAAGGU UGAUG GCAUGCACUAUGC GCG GGUUGUUG	4039
1700	CCGACCUU G AGGCAUAC	1531	GUAGCCU UGAUG GCAUGCACUAUGC GCG AAGGUCCG	4040
1730	UGUUUAU G AGUGGAG	1532	CUCCCAU UGAUG GCAUGCACUAUGC GCG AUUAAACA	4041
1818	AGCACCAU G CAACUUUU	1533	AAAAGUUG UGAUG GCAUGCACUAUGC GCG AUGGUGCU	4042
1835	UCACCUUC G CCUAAUCA	1534	UGAUUAGG UGAUG GCAUGCACUAUGC GCG AGAGGUGA	4043
1883	CAAGCUGU G CCUUGGGU	1535	ACCCAAGG UGAUG GCAUGCACUAUGC GCG ACAGCUUG	4044
1912	UGGACAUU G ACCCGUAU	1536	AUACGGGU UGAUG GCAUGCACUAUGC GCG AAUGUCCA	4045
1959	UCUUUUUU G CCUUCUGA	1537	UCAGAAGG UGAUG GCAUGCACUAUGC GCG AAAAAAGA	4046
1966	UGCCUUUC G ACUUCUUU	1538	AAAGAAGU UGAUG GCAUGCACUAUGC GCG AGAAGGCA	4047
1985	UUCUAUUC G AGAUCUCC	1539	GGAGAUCU UGAUG GCAUGCACUAUGC GCG GAAUAGAA	4048
1996	AUCUCCUC G ACACCGCC	1540	GGCGGUGU UGAUG GCAUGCACUAUGC GCG GAGGAGAU	4049
2002	UCGACACC G CCUCUGCU	1541	AGCAGAGG UGAUG GCAUGCACUAUGC GCG GGUGUCGA	4050
2008	CCGCUCU G CUCUGUAU	1542	AUACAGAG UGAUG GCAUGCACUAUGC GCG AGAGGCGG	4051
2092	GUUGGGGU G AGUUGAUG	1543	CAUCAACU UGAUG GCAUGCACUAUGC GCG ACCCCAAC	4052
2097	GGUGAGUU G AUGAAUCU	1544	AGAUUCAU UGAUG GCAUGCACUAUGC GCG AACUCACC	4053
2100	GAGUUGAU G AAUCUAGC	1545	GCUAGAUU UGAUG GCAUGCACUAUGC GCG AUCAACUC	4054

Table 39

2237	UUUUGGC G AGAAACUG	1546	CAGUUUCU UGAUG GCAUGCACUAUGC GCG GCCAAAA	4055
2251	CUGUUCU G AAUAUUG	1547	CAAAUAU UGAUG GCAUGCACUAUGC GCG AAGAACAG	4056
2282	GUGGAUUC G CACUCCUC	1548	GAGGAGUG UGAUG GCAUGCACUAUGC GCG GAAUCCAC	4057
2293	CUCUCCU G CAUAUAGA	1549	UCUAUAUG UGAUG GCAUGCACUAUGC GCG AGGAGGAG	4058
2311	CACCAAAU G CCCUAUC	1550	GAUAGGGG UGAUG GCAUGCACUAUGC GCG AUUUGGUG	4059
2354	UGUAGAC G AAGAGCA	1551	UGCCUCU UGAUG GCAUGCACUAUGC GCG GUCUAACA	4060
2388	ACUCCUC G CCUCGCAG	1552	CUGGAGG UGAUG GCAUGCACUAUGC GCG GAGGGAGU	4061
2393	CUCGCCUC G CAGACGAA	1553	UUCGUCUG UGAUG GCAUGCACUAUGC GCG GAGGCGAG	4062
2399	UCGCAGAC G AAGGUCUC	1554	GAGACCUU UGAUG GCAUGCACUAUGC GCG GUCUGCGA	4063
2412	UCUCAUUC G CCGCGUCG	1555	CGAGCGG UGAUG GCAUGCACUAUGC GCG GAUUGAGA	4064
2415	CAUUGCC G CGUCGCAG	1556	CUGGACG UGAUG GCAUGCACUAUGC GCG GCGAUUG	4065
2420	GCCGCGUC G CAGAAGAU	1557	AUCUUCUG UGAUG GCAUGCACUAUGC GCG GACGCGC	4066
2514	GGUACCUU G CUUUAUC	1558	GAUUAAG UGAUG GCAUGCACUAUGC GCG AAGGUACC	4067
2549	CUUUUCCU G ACAUUCAU	1559	AUGAAUGU UGAUG GCAUGCACUAUGC GCG AGGAAAG	4068
2560	AUUCAUUU G CAGGAGGA	1560	UCCUCCUG UGAUG GCAUGCACUAUGC GCG AAUUGAU	4069
2576	ACAUGUU G AUAGAUGU	1561	ACAUCUUA UGAUG GCAUGCACUAUGC GCG AACAAUGU	4070
2615	CAGUAAU G AAACAGG	1562	CCUGUUU UGAUG GCAUGCACUAUGC GCG AUUUAUCUG	4071
2641	UUAACUUAU G CCUGCUAG	1563	CUAGCAGG UGAUG GCAUGCACUAUGC GCG AUAGUUA	4072
2645	CUAUGCCU G CUAGGUUU	1564	AAACCUAG UGAUG GCAUGCACUAUGC GCG AGGCAUAG	4073
2677	AAUAUUU G CCUUGA	1565	UCUAAGG UGAUG GCAUGCACUAUGC GCG AAUAUUU	4074
2740	UUCAGAC G CGACAUUA	1566	UUAUGUG UGAUG GCAUGCACUAUGC GCG GUCUGGAA	4075
2742	CCAGACG G ACAUUAU	1567	AAUAUGU UGAUG GCAUGCACUAUGC GCG GCGUCUGG	4076
2804	CACGUAGC G CCUCAUU	1568	AAUGAGG UGAUG GCAUGCACUAUGC GCG GCUACGUG	4077
2814	CUCAUUU G CGGUCAC	1569	GUGACCCG UGAUG GCAUGCACUAUGC GCG AAAUGAG	4078
2875	CAACCCU G AAAGGCA	1570	UGCCUUU UGAUG GCAUGCACUAUGC GCG GAGGUUUG	4079
2928	UCUCCCC G AUCAUCAG	1571	CUGAUGAU UGAUG GCAUGCACUAUGC GCG GGGGAAGA	4080
2946	UGGACCCU G CAUUCAAA	1572	UUUGAAUG UGAUG GCAUGCACUAUGC GCG AGGJUCCA	4081
2990	CUCAACCC G CACAAGGA	1573	UCCUUGUG UGAUG GCAUGCACUAUGC GCG GGGUUGAG	4082
3012	GGCGGAC G CCAACAAG	1574	CUUUGUG UGAUG GCAUGCACUAUGC GCG GUCCGGCC	4083
3090	GCCUUCAC G CUCAGGGC	1575	GCCUGAG UGAUG GCAUGCACUAUGC GCG GUGAGGCG	4084
3113	ACAACUGU G CCAGCAGC	1576	GCUGCUGG UGAUG GCAUGCACUAUGC GCG ACAGUUGU	4085
3132	CUCUCCU G CCUCCACC	1577	GGUGGAGG UGAUG GCAUGCACUAUGC GCG AGGAGGAG	4086
51	AGGGCCCU G UACUUUCC	1578	GGAAAGUA UGAUG GCAUGCACUAUGC GCG AGGGCCCU	4087
106	AGAAUACU G UCUCUGCC	1579	GGCAGAGA UGAUG GCAUGCACUAUGC GCG AGUAUUCU	4088

Table 39

148	GGGACCCU G UACCGAAC	1580	GUUCGUUA UGAUG GCAUGCACUAUGC GCG AGGGUCCC	4089
198	CUGCUGU G UACAGGC	1581	GCCGUAA UGAUG GCAUGCACUAUGC GCG ACGAGCAG	4090
219	UUUUUUU G UUGACAAA	1582	UUUGCAA UGAUG GCAUGCACUAUGC GCG AAGAAAAA	4091
297	ACACCCG G UGUCUUGG	1583	CCAAGACA UGAUG GCAUGCACUAUGC GCG ACGGGUGU	4092
299	ACCCGUG G UCUUGGCC	1584	GGCCAAGA UGAUG GCAUGCACUAUGC GCG ACACGGGU	4093
347	ACCAACCU G UUGUCCUC	1585	GAGGACAA UGAUG GCAUGCACUAUGC GCG AGGUUGGU	4094
350	AACCUGU G UCCUCCAA	1586	UUGGAGGA UGAUG GCAUGCACUAUGC GCG AACAGGUU	4095
362	UCCAUAU G UCCUGGUU	1587	AACCAGGA UGAUG GCAUGCACUAUGC GCG AAUUGGA	4096
381	CGCUGGAU G UGUCUGCG	1588	CGCAGACA UGAUG GCAUGCACUAUGC GCG AUCCAGCG	4097
383	CUGGAUGU G UCUGCGGC	1589	GCCGACGA UGAUG GCAUGCACUAUGC GCG ACAUCCAG	4098
438	AUCUUCU G UUGGUUCU	1590	AGAACCAA UGAUG GCAUGCACUAUGC GCG AAGAAGAU	4099
465	CAAGGUU G UUGCCCGU	1591	ACGGCAA UGAUG GCAUGCACUAUGC GCG AUACCUUG	4100
476	GCCGUAU G UCCUUA	1592	UUAGAGGA UGAUG GCAUGCACUAUGC GCG AAACGGGC	4101
555	ACCUCUAU G UUUCCUUC	1593	GAGGAAA UGAUG GCAUGCACUAUGC GCG AUAGAGGU	4102
566	UCCUUAU G UUGCUGUA	1594	UACAGCAA UGAUG GCAUGCACUAUGC GCG AUGAGGGA	4103
572	AUGUUGU G UACAAAAC	1595	GUUUUGUA UGAUG GCAUGCACUAUGC GCG AGCAACAU	4104
602	CUGACCU G UAUUCCCA	1596	UGGGAUA UGAUG GCAUGCACUAUGC GCG AGGUGCAG	4105
694	UGCCAUAU G UUCAGUGG	1597	CCACUGAA UGAUG GCAUGCACUAUGC GCG AAUUGGCA	4106
724	CCCCACU G UCUGGCUU	1598	AAGCCAGA UGAUG GCAUGCACUAUGC GCG AGUGGGGG	4107
750	UGGAUGAU G UGGUUUUG	1599	CAAAACCA UGAUG GCAUGCACUAUGC GCG AUCAUCCA	4108
771	CCAAGUCU G UACAACAU	1600	AUGUUUGA UGAUG GCAUGCACUAUGC GCG AGACTUUG	4109
801	AUGCGCU G UUAACCAU	1601	AUUGGUA UGAUG GCAUGCACUAUGC GCG AGCGGCAU	4110
818	UUUUAUU G UCUUUGGG	1602	CCCAAGA UGAUG GCAUGCACUAUGC GCG AAAAGAAA	4111
888	UGGGAUAU G UAAUUGGG	1603	CCCAUAUA UGAUG GCAUGCACUAUGC GCG AUAUCCCA	4112
927	AACAUAU G UACAAAAA	1604	UUUUUGUA UGAUG GCAUGCACUAUGC GCG AAUAUGUU	4113
944	AUCAAUAU G UGUUUUAG	1605	CUAAAACA UGAUG GCAUGCACUAUGC GCG AUUUUGAU	4114
946	CAAAUAU G UUUUAAGA	1606	UCCUAAA UGAUG GCAUGCACUAUGC GCG ACAUUUUG	4115
963	AACUCCU G UAAACAGG	1607	CCUGUUUA UGAUG GCAUGCACUAUGC GCG AGGAAGUU	4116
991	GAAAGUAU G UCAACGAA	1608	UUUGUUGA UGAUG GCAUGCACUAUGC GCG AUACUUUC	4117
1002	AACGAUAU G UGGGUCUU	1609	AAGACCCA UGAUG GCAUGCACUAUGC GCG AAUUGGUU	4118
1039	CAGCAUAU G UGGAUAU	1610	AAUAUCCA UGAUG GCAUGCACUAUGC GCG AUUGCGUG	4119
1137	AACAGUAU G UGAACCUU	1611	AAGUUUCA UGAUG GCAUGCACUAUGC GCG AUACUGUU	4120
1184	UGCAAGU G UUUGCUGA	1612	UCAGCAAA UGAUG GCAUGCACUAUGC GCG ACUUGGCA	4121
1251	GAACCUU G UGUCCUCC	1613	AGGAGACA UGAUG GCAUGCACUAUGC GCG AAAGGUUC	4122

Table 39

1253	ACCUJUGU G UCUCCUCU	1614	AGAGGAGA UGAUG GCAUGCACUAUGC GCG ACAAAGGU	4123
1294	AGCCGCUU G UUUUGCUC	1615	GAGCAAAA UGAUG GCAUGCACUAUGC GCG AAGCGGCU	4124
1344	ACAAUUCU G UCGUGCUC	1616	GAGCAGCA UGAUG GCAUGCACUAUGC GCG AGAAUUGU	4125
1390	GUAGGCU G UGCUGCCA	1617	UGGCAGCA UGAUG GCAUGCACUAUGC GCG AGCCUAGC	4126
1425	CGUCCUUU G UUUACGUC	1618	GACGUAAA UGAUG GCAUGCACUAUGC GCG AAAGGACG	4127
1508	CGCCUAUU G UACCGACC	1619	GGUCGUUA UGAUG GCAUGCACUAUGC GCG AAUAGGCG	4128
1557	CCCCGUCU G UGCCUUCU	1620	AGAGGCA UGAUG GCAUGCACUAUGC GCG AGACGGGG	4129
1581	CGGACCGU G UGCACUUC	1621	GAAGUGCA UGAUG GCAUGCACUAUGC GCG ACGGUCCG	4130
1684	UCAGCAAU G UCAACGAC	1622	GUCGUUGA UGAUG GCAUGCACUAUGC GCG AUUGCUGA	4131
1719	CAAGACU G UGUGUUUA	1623	UAAACACA UGAUG GCAUGCACUAUGC GCG AGUCUUUG	4132
1721	AAGACUGU G UGUUUAAU	1624	AUUAACA UGAUG GCAUGCACUAUGC GCG ACAGUCUU	4133
1723	GACUGUGU G UUUUAUGA	1625	UCAUUAAA UGAUG GCAUGCACUAUGC GCG ACACAGUC	4134
1772	AGGUCUUU G UACUAGGA	1626	UCCUAGUA UGAUG GCAUGCACUAUGC GCG AAAGACCU	4135
1785	AGGAGGCU G UAGGCAUA	1627	UUGCCUA UGAUG GCAUGCACUAUGC GCG AGCCUCCU	4136
1801	AAAUUGGU G UGUUCACC	1628	GGUGAACA UGAUG GCAUGCACUAUGC GCG ACCAAUUU	4137
1803	AUUGGUGU G UUCACCAG	1629	CUGUGAA UGAUG GCAUGCACUAUGC GCG ACACCAAU	4138
1850	CAUCUACU G UUCAUGUC	1630	GACAUGAA UGAUG GCAUGCACUAUGC GCG AUGAGAUG	4139
1856	AUGUACAU G UCCUACUG	1631	CAGUAGGA UGAUG GCAUGCACUAUGC GCG AUGAACAU	4140
1864	GUCCUACU G UUCAAGCC	1632	GGUUGAA UGAUG GCAUGCACUAUGC GCG AGUAGGAC	4141
1881	UCCRAGCU G UGCCUUGG	1633	CCRAGGCA UGAUG GCAUGCACUAUGC GCG AGCUUGGA	4142
1939	GAGCUUCU G UGGAGUUA	1634	UAAUCUCA UGAUG GCAUGCACUAUGC GCG AGAAGCUC	4143
2013	UCUGUCU G UAUCGGGG	1635	CCCCGAUA UGAUG GCAUGCACUAUGC GCG AGAGCAGA	4144
2045	GGAACAUU G UUCACCUC	1636	GAGUGAA UGAUG GCAUGCACUAUGC GCG AAUGUJCC	4145
2082	GUUAUUCU G UGUJGGGG	1637	CCCCRACA UGAUG GCAUGCACUAUGC GCG AGAAUJAGC	4146
2084	UAUUCUGU G UUGGGGUG	1638	CACCCCAA UGAUG GCAUGCACUAUGC GCG ACAGAAUA	4147
2167	UCAGCUAU G UCAACGJU	1639	AACGUJGA UGAUG GCAUGCACUAUGC GCG AUAGCTGA	4148
2205	CAACUAUU G UGGUJUUA	1640	UGAAACCA UGAUG GCAUGCACUAUGC GCG AAUAGUJUG	4149
2222	CAUJUCCU G UCUUACUJ	1641	AAGUAAGA UGAUG GCAUGCACUAUGC GCG AGGAAUUG	4150
2245	GAGAAACU G UUCUUGAA	1642	UUCAAAGNA UGAUG GCAUGCACUAUGC GCG AGUJUUCUC	4151
2262	UAUJUGGU G UCUJUUGG	1643	CCAAAAGA UGAUG GCAUGCACUAUGC GCG ACCAAUAU	4152
2274	UUUGGAGU G UGGAUUCG	1644	CGAAUCCA UGAUG GCAUGCACUAUGC GCG ACUCCAAA	4153
2344	AAACUACU G UUGUJAGA	1645	UCUAAACNA UGAUG GCAUGCACUAUGC GCG AGUAGUJU	4154
2347	CUACUGUU G UJAGACGA	1646	UCGUCUNA UGAUG GCAUGCACUAUGC GCG AACAGUAG	4155
2450	AUCUCAAU G UJAGUAUU	1647	AAUACUAA UGAUG GCAUGCACUAUGC GCG AUTGAGAU	4156

Table 39

2573	AGGACAUU G UGAUAGA	1648	UCUAUCAA UGAUG GCAUGCACUAUGC GCG AAUGUCCU	4157
2583	UGAUAGAU G UAAAGCAAU	1649	AUUGCUUA UGAUG GCAUGCACUAUGC GCG AUCUAUCA	4158
2594	AGCAAUUU G UGGGGCCC	1650	GGGCCCCA UGAUG GCAUGCACUAUGC GCG AAUUGCU	4159
2663	AUCCCAAU G UUAUAAA	1651	UUUAGUAA UGAUG GCAUGCACUAUGC GCG AUUGGGAU	4160
2717	CAGAGUAA G UAGUUAAU	1652	AUUAACTA UGAUG GCAUGCACUAUGC GCG AUACUCUG	4161
2901	AUCUUUCU G UCCCCAAU	1653	AUUGGGGA UGAUG GCAUGCACUAUGC GCG AGAAAGAU	4162
3071	GGGGACU G UUGGGGUG	1654	CACCCCAA UGAUG GCAUGCACUAUGC GCG AGUCCCCC	4163
3111	UCACAAAU G UGCCAGCA	1655	UGCUGGCA UGAUG GCAUGCACUAUGC GCG AGUUGUGA	4164

Input Sequence = AF100308. Cut Site = YG/M or UG/U.
 Stem Length = 8. Core Sequence = UGAUG GCAUGCACUAUGC GCG
 AF100308 (Hepatitis B virus strain 2-18, 3215 bp)

Table 40

Table 40: Human HBV Zinzyme Ribozyme and Substrate Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
61	ACUUUCCU G CUGUGGC	1448	GCCACCAG GCcgaagGCGaGuCaaGGGuCu AGGAAAAGU	4165
94	UGAGCCCU G CUCAGAAU	1450	AUUCUGAG GCcgaagGCGaGuCaaGGGuCu AGGGCUCA	4166
112	CUGUCUCU G CCAUAUCG	1451	CGAUUUGG GCcgaagGCGaGuCaaGGGuCu AGAGACAG	4167
169	AGAACAU G CAUCAGGA	1454	UCCUGAUG GCcgaagGCGaGuCaaGGGuCu GAUGUUCU	4168
192	GGACCCCU G CUCGUGUU	1455	AACACGAG GCcgaagGCGaGuCaaGGGuCu AGGGGUCC	4169
315	CAAAAUUC G CAGUCCCA	1457	UGGACUG GCcgaagGCGaGuCaaGGGuCu GAAUUUUG	4170
374	UGGUUAUC G CUGGAUGU	1458	ACAUCAG GCcgaagGCGaGuCaaGGGuCu GAUAACCA	4171
387	AUGUGUCU G CGGCGUUU	1459	AAACGCCG GCcgaagGCGaGuCaaGGGuCu AGACACAU	4172
410	CUUCCUCU G CAUCCUCC	1460	GCAGGAG GCcgaagGCGaGuCaaGGGuCu AGAGGAAG	4173
417	UGCAUCCU G CUGCUAUG	1461	CAUAGCAG GCcgaagGCGaGuCaaGGGuCu AGGAUGCA	4174
420	AUCCUGCU G CUAUGCCU	1462	AGGCAUAG GCcgaagGCGaGuCaaGGGuCu AGCAGGAU	4175
425	GUCGUAU G CCUCAUCU	1463	AGAUAGG GCcgaagGCGaGuCaaGGGuCu AUAGCAGC	4176
468	GGUAUGUU G CCCGUUUG	1464	CAACGGG GCcgaagGCGaGuCaaGGGuCu AACAUACC	4177
518	CGACCAU G CAAAACCU	1465	AGGUUUUG GCcgaagGCGaGuCaaGGGuCu AUGGUCCG	4178
527	CAAAACCU G CACACUC	1466	GAGUUUG GCcgaagGCGaGuCaaGGGuCu AGGUUUUG	4179
538	CAACUCCU G CUCAAGGA	1467	UCCUUAG GCcgaagGCGaGuCaaGGGuCu AGGAGUUG	4180
569	CUCAUGUU G CUGUACAA	1468	UUGUACAG GCcgaagGCGaGuCaaGGGuCu AACAUAG	4181
596	CGGAAACU G CACCUGUA	1469	UACAGGUG GCcgaagGCGaGuCaaGGGuCu AGUUUCCG	4182
631	GGGUUUC G CAAAUAAC	1470	GUUUUUUG GCcgaagGCGaGuCaaGGGuCu GAAAGCCC	4183
687	UUACUAGU G CCAUUUUG	1471	ACAAUUG GCcgaagGCGaGuCaaGGGuCu ACUAGUAA	4184
795	CCUUUAU G CCGCUGUU	1474	AACAGCG GCcgaagGCGaGuCaaGGGuCu AUAAAGGG	4185
798	UUUAUGCC G CUGUUAAC	1475	GGUAAACG GCcgaagGCGaGuCaaGGGuCu GGCAUAAA	4186
911	GGCACAUU G CCACAGGA	1476	UCCUGUGG GCcgaagGCGaGuCaaGGGuCu AAUGUGCC	4187
1020	UGGGUUU G CCGCCCUU	1479	AGGGCGG GCcgaagGCGaGuCaaGGGuCu AAACCCCA	4188
1023	GGUUUGCC G CCCUUUUC	1480	GAAAGGG GCcgaagGCGaGuCaaGGGuCu GGCAAAAC	4189
1034	CCUUUCAC G CAAUGUGG	1481	CCACAUUG GCcgaagGCGaGuCaaGGGuCu GUGAAAGG	4190
1050	GAUAUUCU G CUUUAAUG	1482	CAUUAAG GCcgaagGCGaGuCaaGGGuCu AGAAUADC	4191
1058	GUUUAAU G CCUUUAUA	1483	UAUUAAG GCcgaagGCGaGuCaaGGGuCu AUUAAGC	4192
1068	CUUAUAU G CAUGCAUA	1484	UAUGCAUG GCcgaagGCGaGuCaaGGGuCu AUUAAGG	4193
1072	AUAUGCAU G CAUACAAG	1485	CUUGUAUG GCcgaagGCGaGuCaaGGGuCu AUGCAUAU	4194

Table 40

1103	ACUUCUC G CCAACUUA	1486	UAAGUUG GCCgaaagGCGaGuCaaGGuCu GAGAAAGU	4195
1155	ACCCCGU G CUCGGCAA	1488	UUGCCGAG GCCgaaagGCGaGuCaaGGuCu AACGGGU	4196
1177	UGGUCUAU G CCAAGUGU	1489	ACACUUG GCCgaaagGCGaGuCaaGGuCu AUAGACCA	4197
1188	AAGUGUUU G CUGACGCA	1490	UUCGUCAG GCCgaaagGCGaGuCaaGGuCu AAACACUU	4198
1194	UUGCUGAC G CAACCCCC	1492	GGGGUUG GCCgaaagGCGaGuCaaGGuCu GUCAGCAA	4199
1234	CAUCAGC G CAUGCUG	1493	CACGCAUG GCCgaaagGCGaGuCaaGGuCu GCUGAUGG	4200
1238	CAGCGAU G CGUGGAAC	1494	GUUCCACG GCCgaaagGCGaGuCaaGGuCu AUGCGCUG	4201
1262	UCUCUCU G CCGAUCCA	1495	UGGAUCGG GCCgaaagGCGaGuCaaGGuCu AGAGGAGA	4202
1275	UCCAUACC G CGGAACUC	1497	GAGUUCGG GCCgaaagGCGaGuCaaGGuCu GGUUAUGGA	4203
1290	UCCUAGCC G CUUGUUUU	1498	AAACACAG GCCgaaagGCGaGuCaaGGuCu GGCUAGGA	4204
1299	CUUGUUUU G CUCGCAGC	1499	GCUGCGAG GCCgaaagGCGaGuCaaGGuCu AAAACAAG	4205
1303	UUUGUCU G CAGCAGGU	1500	ACCUGCUG GCCgaaagGCGaGuCaaGGuCu GAGCAAAA	4206
1349	UCUGUCGU G CUCUCCCG	1502	CGGGAGAG GCCgaaagGCGaGuCaaGGuCu ACGACAGA	4207
1357	GCUCUCC G CAAAUUA	1503	UAUAUUUG GCCgaaagGCGaGuCaaGGuCu GGGAGAGC	4208
1382	CAUGGCU G CUAGGCU	1504	CAGCCUAG GCCgaaagGCGaGuCaaGGuCu AGCCAUGG	4209
1392	UAGGUCU G CUGCCAAC	1505	GUUGGAG GCCgaaagGCGaGuCaaGGuCu ACAGCCUA	4210
1395	GCUGUCU G CCAACUGG	1506	CCAGUUUG GCCgaaagGCGaGuCaaGGuCu AGCACAGC	4211
1411	GAUCCUAC G CGGGACGU	1507	ACGUCCCG GCCgaaagGCGaGuCaaGGuCu GUAGGAUC	4212
1442	CCGUGGC G CUGAAUCC	1508	GGAUUCAG GCCgaaagGCGaGuCaaGGuCu GCCGACGG	4213
1452	UGAAUCCC G CGGACGAC	1510	GUCGUCCG GCCgaaagGCGaGuCaaGGuCu GGGAUUCA	4214
1474	CCGGGGCC G CUUGGGGC	1512	GCCCCAAG GCCgaaagGCGaGuCaaGGuCu GGGCCCGG	4215
1489	GCUCUACC G CCCGCUUC	1513	GAAGCGGG GCCgaaagGCGaGuCaaGGuCu GGUAGAGC	4216
1493	UACGCCC G CUUCUCCG	1514	CGGAGNAG GCCgaaagGCGaGuCaaGGuCu GGGCGGUA	4217
1501	GCUUCUCC G CCUAUUGU	1515	ACAAUAGG GCCgaaagGCGaGuCaaGGuCu GGAGAAGC	4218
1528	CACGGGC G CACUUCUC	1517	GAGAGGUG GCCgaaagGCGaGuCaaGGuCu GCGCCGUG	4219
1542	CUUUUAC G CGGACUCC	1518	GGAGUCCG GCCgaaagGCGaGuCaaGGuCu GUAAAGAG	4220
1559	CCGUCUGU G CCUUCUCA	1519	UGAGAAGG GCCgaaagGCGaGuCaaGGuCu ACAGACGG	4221
1571	UCUCAUCU G CCGGACCG	1520	CGGUCCGG GCCgaaagGCGaGuCaaGGuCu AGAUGAGA	4222
1583	GACCGUGU G CACUUCGC	1521	GCGAAGUG GCCgaaagGCGaGuCaaGGuCu ACACGGUC	4223
1590	UGCACUUC G CUUCACCU	1522	AGGUAAG GCCgaaagGCGaGuCaaGGuCu GAAGUGCA	4224
1601	UCACUCU G CACGUCGC	1523	GCGACGUG GCCgaaagGCGaGuCaaGGuCu AGAGGUGA	4225
1608	UGCACGUC G CAUGGAGA	1524	UCUCCAUU GCCgaaagGCGaGuCaaGGuCu GACGUGCA	4226
1628	CCUGAAC G CCCACAGG	1526	CCUGUGGG GCCgaaagGCGaGuCaaGGuCu GUTACACGG	4227
1642	AGGAACCU G CCCAAGGU	1527	ACCUUGGG GCCgaaagGCGaGuCaaGGuCu AGGUUCCU	4228

Table 40

1654	AAGGUCUU	G	CAUAAGAG	1528	CUCUUAUG	GCcgaagGCGaGuCaaGGuCu	AAGACCUU	4229
1818	AGCACCAU	G	CAACUJU	1533	AAAAGUUG	GCcgaagGCGaGuCaaGGuCu	AUGGUGCU	4230
1835	UCACCUCU	G	CCUAUUA	1534	UGAUUAGG	GCcgaagGCGaGuCaaGGuCu	AGAGGUGA	4231
1883	CAAGCUGU	G	CCUUGGGU	1535	ACCCAAGG	GCcgaagGCGaGuCaaGGuCu	ACAGCUUG	4232
1959	UCUJUJU	G	CCUUCUGA	1537	UCAGAAGG	GCcgaagGCGaGuCaaGGuCu	AAAAAGA	4233
2002	UCGACACC	G	CCUCUGCU	1541	AGCAGAGG	GCcgaagGCGaGuCaaGGuCu	GGUGUCGA	4234
2008	CCGCCUCU	G	CUCUGUAU	1542	AUACAGAG	GCcgaagGCGaGuCaaGGuCu	AGAGGGGG	4235
2282	GUGGAUUC	G	CACUCCUC	1548	GAGGAGUG	GCcgaagGCGaGuCaaGGuCu	GAUCCAC	4236
2293	CUCUCCU	G	CAUAUAGA	1549	UCUAUAUG	GCcgaagGCGaGuCaaGGuCu	AGGAGGAG	4237
2311	CACCAAAU	G	CCCCUAUC	1550	GAUAGGGG	GCcgaagGCGaGuCaaGGuCu	AUTUGGUG	4238
2388	ACUCCUC	G	CCUCGCAG	1552	CUGCGAGG	GCcgaagGCGaGuCaaGGuCu	GAGGGAGU	4239
2393	CUCGCCUC	G	CAGACGAA	1553	UUCGUCUG	GCcgaagGCGaGuCaaGGuCu	GAGGCGAG	4240
2412	UCUCAUUC	G	CCGCGUCG	1555	CGACGCGG	GCcgaagGCGaGuCaaGGuCu	GAUUGAGA	4241
2415	CAUUGCC	G	CGUCGCAG	1556	CUGCGACG	GCcgaagGCGaGuCaaGGuCu	GGCGAUUG	4242
2420	GCCGCGUC	G	CAGAAGAU	1557	AUCUUCUG	GCcgaagGCGaGuCaaGGuCu	GACGCGGC	4243
2514	GGUACCUU	G	CUUUAUUC	1558	GAUUAAG	GCcgaagGCGaGuCaaGGuCu	AAGGUACC	4244
2560	AUUCAUUU	G	CAGGAGGA	1560	UCCUUCUG	GCcgaagGCGaGuCaaGGuCu	AAUGAAU	4245
2641	UUAACUUA	G	CCUGCUAG	1563	CUAGCAGG	GCcgaagGCGaGuCaaGGuCu	AUAGUUA	4246
2645	CUAUGCCU	G	CUAGGUUU	1564	AAACCUAG	GCcgaagGCGaGuCaaGGuCu	AGGCAUAG	4247
2677	AAUAUUUU	G	CCCUUAGA	1565	UCUAAGGG	GCcgaagGCGaGuCaaGGuCu	AAUAUUU	4248
2740	UUCCAGAC	G	CGACAUUA	1566	UAAUGUCG	GCcgaagGCGaGuCaaGGuCu	GUCUGGAA	4249
2804	CACGUAGC	G	CCUCAUUU	1568	AAUAGAGG	GCcgaagGCGaGuCaaGGuCu	GUUACGUG	4250
2814	CUCAUUUU	G	CGGGUCAC	1569	GUGACCCG	GCcgaagGCGaGuCaaGGuCu	AAAUAGAG	4251
2946	UGGACCCU	G	CAUUCNAA	1572	UUUGAAUG	GCcgaagGCGaGuCaaGGuCu	AGGGUCCA	4252
2990	CUCAACCC	G	CAUUCNAA	1573	UCCUUGUG	GCcgaagGCGaGuCaaGGuCu	GGGUUGAG	4253
3012	GGCGGAC	G	CCAACNAG	1574	CUUGUUUG	GCcgaagGCGaGuCaaGGuCu	GUCCGGCC	4254
3090	GCCUUCAC	G	CCACGAGC	1575	GCCUUGAG	GCcgaagGCGaGuCaaGGuCu	GUAGGGGC	4255
3113	ACACUGU	G	CCAGCAGC	1576	GUUGUUGG	GCcgaagGCGaGuCaaGGuCu	ACAGUUUG	4256
3132	CUCCUCCU	G	CCUCCACC	1577	GGUGGAGG	GCcgaagGCGaGuCaaGGuCu	AGGAGGAG	4257
51	AGGGCCCU	G	UACUJUCC	1578	GGAAAGUA	GCcgaagGCGaGuCaaGGuCu	AGGGCCCU	4258
106	AGAAUACU	G	UCUCUGCC	1579	GGCAGAGA	GCcgaagGCGaGuCaaGGuCu	AGUAUUCU	4259
148	GGGACCCU	G	UACCGAAC	1580	GUUCGUUA	GCcgaagGCGaGuCaaGGuCu	AGGUGUCC	4260
198	CUGCUCGU	G	UUACAGGC	1581	GCCUGUAA	GCcgaagGCGaGuCaaGGuCu	ACGAGCAG	4261
219	UUUUUCUU	G	UUGACAAA	1582	UUUGUCAA	GCcgaagGCGaGuCaaGGuCu	AAGAAAAA	4262

Table 40

297	ACACCCGU	G	UGUCUUGG	1583	CCAAGACA	GCcgaagGCGaGuCaaGGuCu	ACGGUGU	4263
299	ACCCGUGU	G	UCUUGGCC	1584	GGCCAAGA	GCcgaagGCGaGuCaaGGuCu	ACACGGGU	4264
347	ACCAACCU	G	UUGUCCUC	1585	GAGGACAA	GCcgaagGCGaGuCaaGGuCu	AGGUUGGU	4265
350	AACCUGUU	G	UCCUCCAA	1586	UUGGAGGA	GCcgaagGCGaGuCaaGGuCu	AACAGGUU	4266
362	UCCAUAUU	G	UCCUGGUU	1587	AACCAGGA	GCcgaagGCGaGuCaaGGuCu	AAAUUGGA	4267
381	CGCUGGAU	G	UGUCUGCG	1588	CGCAGACA	GCcgaagGCGaGuCaaGGuCu	AUCCAGCG	4268
383	CUGGAUGU	G	UCUGCGGC	1589	GCCGCAGA	GCcgaagGCGaGuCaaGGuCu	ACAUCCAG	4269
438	AUCUUCUU	G	UUGGUUCU	1590	AGAACCAG	GCcgaagGCGaGuCaaGGuCu	AAGAAGAU	4270
465	CAAGGUUU	G	UUGCCCGU	1591	ACGGGCAA	GCcgaagGCGaGuCaaGGuCu	AUACCUUG	4271
476	GCCCGUUU	G	UCCUCUAA	1592	UUAGAGGA	GCcgaagGCGaGuCaaGGuCu	AAACGGGC	4272
555	ACCUUAUU	G	UUUCCUCU	1593	GAGGGAAG	GCcgaagGCGaGuCaaGGuCu	AUAGAGGU	4273
566	UCCUUAUU	G	UUGCUGUA	1594	UACAGCAA	GCcgaagGCGaGuCaaGGuCu	AUGAGGGA	4274
572	AUGUUGCU	G	UACAAAC	1595	GUUUUGUA	GCcgaagGCGaGuCaaGGuCu	AGCAACAU	4275
602	UGCCAUUU	G	UAUUCCCA	1596	UGGAAUA	GCcgaagGCGaGuCaaGGuCu	AGGUGCAG	4276
694	UGCCAUUU	G	UAUUCCCA	1597	CCACUGAA	GCcgaagGCGaGuCaaGGuCu	AAAUUGGA	4277
724	CCCCACUU	G	UCUGGCUU	1598	AAGCCAGA	GCcgaagGCGaGuCaaGGuCu	AGUGGGGG	4278
750	UGGAUGAU	G	UGGUUUUG	1599	CAAAACCA	GCcgaagGCGaGuCaaGGuCu	AUCAUCCA	4279
771	CCAAGUCU	G	UACAACAU	1600	AUUUGUA	GCcgaagGCGaGuCaaGGuCu	AGACUUGG	4280
801	AUGCCGCU	G	UUACCAAU	1601	AUUGGUA	GCcgaagGCGaGuCaaGGuCu	AGCGGCAU	4281
818	UGGUUUUU	G	UCUUUGGG	1602	CCCAAGA	GCcgaagGCGaGuCaaGGuCu	AAAAGAAA	4282
888	UGGAUAUU	G	UAUUUGGG	1603	CCCAUUA	GCcgaagGCGaGuCaaGGuCu	AUAUCCCA	4283
927	AACAUAUU	G	UACAAAA	1604	UUUUUGUA	GCcgaagGCGaGuCaaGGuCu	AAUAUGUU	4284
944	AUCAAUUU	G	UGUUUAG	1605	CUAAAA	GCcgaagGCGaGuCaaGGuCu	AUUUUGAU	4285
946	CAAAAUUU	G	UUUUAGGA	1606	UCCUAAA	GCcgaagGCGaGuCaaGGuCu	ACAUUUUG	4286
963	AACUUCUU	G	UAAACAGG	1607	CTUGUUA	GCcgaagGCGaGuCaaGGuCu	AGGAAGUU	4287
991	GAAGUAUU	G	UCAACGAA	1608	UUCGUUA	GCcgaagGCGaGuCaaGGuCu	AUACUUUC	4288
1002	AACGAUUU	G	UGGGUCUU	1609	AAGACCCA	GCcgaagGCGaGuCaaGGuCu	AAUUCGUU	4289
1039	CACGAUUU	G	UGGAUAUU	1610	AAUAUCCA	GCcgaagGCGaGuCaaGGuCu	AUUUCGUU	4290
1137	AACAGUAU	G	UGAACCUU	1611	AAGGUUA	GCcgaagGCGaGuCaaGGuCu	AUACUGUU	4291
1184	UGCAAGUU	G	UUUGCUGA	1612	UCAGCAAA	GCcgaagGCGaGuCaaGGuCu	ACUUGGCA	4292
1251	GAACUUUU	G	UGUCUCCU	1613	AGGAGACA	GCcgaagGCGaGuCaaGGuCu	AAAGGUUC	4293
1253	ACCUUUGU	G	UCUCCUCU	1614	AGAGGAGA	GCcgaagGCGaGuCaaGGuCu	ACAAAGGU	4294
1294	AGCGCUUU	G	UUUUGCUC	1615	GAGCAAAA	GCcgaagGCGaGuCaaGGuCu	AAGCGGCU	4295
1344	ACAAUUCU	G	UCUGUCUC	1616	GAGCACGA	GCcgaagGCGaGuCaaGGuCu	AGAAUUGU	4296

Table 40

1390	GUUAGGCU G UGUUGCCA	1617	UGGCAGCA GCGaaagCGGaGuCaaGGuCu AGCCUAGC	4297
1425	CGUCCUUU G UUUAGGUC	1618	GACGUAAA GCGaaagCGGaGuCaaGGuCu AAAGGACG	4298
1508	CGCCUAUU G UACCGACC	1619	GGUCGGUA GCGaaagCGGaGuCaaGGuCu AAUAGGCG	4299
1557	CCCGUCU G UGCUUUCU	1620	AGAAGCA GCGaaagCGGaGuCaaGGuCu AGACGGGG	4300
1581	CGGACCGU G UGCACUUC	1621	GAAGUGCA GCGaaagCGGaGuCaaGGuCu ACGGUCCG	4301
1684	UCAGCAAU G UCAACGAC	1622	GUCGUUGA GCGaaagCGGaGuCaaGGuCu AUUGCTGA	4302
1719	CAAAGACU G UGUUUUA	1623	UAAACACA GCGaaagCGGaGuCaaGGuCu AGUCUUUG	4303
1721	AAGACUGU G UGUUUAAU	1624	AUTAAACA GCGaaagCGGaGuCaaGGuCu ACAGUCUU	4304
1723	GACUGUGU G UUUAAUGA	1625	UCAUUAUA GCGaaagCGGaGuCaaGGuCu ACACAGUC	4305
1772	AGGUCUUU G UACUAGGA	1626	UCCUAGUA GCGaaagCGGaGuCaaGGuCu AAAGACCU	4306
1785	AGGAGGCU G UAGGCAUA	1627	UAUGCCUA GCGaaagCGGaGuCaaGGuCu AGCCUCCU	4307
1801	AAAUUGGU G UGUUACCC	1628	GGUGAACA GCGaaagCGGaGuCaaGGuCu ACCAAUUU	4308
1803	AUUGGUGU G UUCACCCG	1629	CUGUGUAA GCGaaagCGGaGuCaaGGuCu ACACCAAU	4309
1850	CAUCUCAU G UUCAUGUC	1630	GACAUCAA GCGaaagCGGaGuCaaGGuCu AUGAGAUG	4310
1856	AUGUUCAU G UCCUACUG	1631	CAGUAGGA GCGaaagCGGaGuCaaGGuCu AUGAACAU	4311
1864	GUCCUACU G UUCAAGCC	1632	GGUUUCAA GCGaaagCGGaGuCaaGGuCu AGUAGGAC	4312
1881	UCCAAGCU G UGCUUUGG	1633	CCAAGGCA GCGaaagCGGaGuCaaGGuCu AGCUUGGA	4313
1939	GAGCUUCU G UGGAGUUA	1634	UAUCUCCA GCGaaagCGGaGuCaaGGuCu AGAAGCUC	4314
2013	UCUGUCU G UAUCCGGG	1635	CCCCAUA GCGaaagCGGaGuCaaGGuCu AGAGCAGA	4315
2045	GGACAUU G UUCACCTC	1636	GAGGUGAA GCGaaagCGGaGuCaaGGuCu AAUGUUCC	4316
2082	GUUAUUCU G UGUUGGGG	1637	CCCACAA GCGaaagCGGaGuCaaGGuCu AGAAUAGC	4317
2084	UAUUCUGU G UUGGGGUG	1638	CACCCAA GCGaaagCGGaGuCaaGGuCu ACAGAAUA	4318
2167	UCAGCUAU G UCAACGUU	1639	AACGUUGA GCGaaagCGGaGuCaaGGuCu AUAGCTGA	4319
2205	CAACUAUU G UGGUUUCA	1640	UGAAACCA GCGaaagCGGaGuCaaGGuCu AGGAAUUG	4320
2222	CAUUUCCU G UCUUACUU	1641	AAGUAAGA GCGaaagCGGaGuCaaGGuCu AGUUUUCU	4321
2245	GAGAAACU G UUCUUGAA	1642	UUCAAGAA GCGaaagCGGaGuCaaGGuCu AGUUUUCU	4322
2262	UAUUUGGU G UCUUUUGG	1643	CCAAAGA GCGaaagCGGaGuCaaGGuCu ACCAAUAU	4323
2274	UUUGGAGU G UGGAUUCG	1644	CGAAUCCA GCGaaagCGGaGuCaaGGuCu ACUCCAAA	4324
2344	AAACUACU G UUGUUAGA	1645	UCUAACAA GCGaaagCGGaGuCaaGGuCu AGUAGUUU	4325
2347	CUACUGUU G UUAGACGA	1646	UCGUUCAA GCGaaagCGGaGuCaaGGuCu AACAGUAG	4326
2450	AUCUCAAU G UUAUAUUU	1647	AAUACUAA GCGaaagCGGaGuCaaGGuCu AUUGAGAU	4327
2573	AGGACAUU G UUGAUAGA	1648	UCUAUCAA GCGaaagCGGaGuCaaGGuCu AAUGUCCU	4328
2583	UGAUAGAU G UAAGCAAU	1649	AUUGCUUA GCGaaagCGGaGuCaaGGuCu AUCUAUCA	4329
2594	AGCAAUUU G UGGGGCCC	1650	GGGCCCCA GCGaaagCGGaGuCaaGGuCu AAUUGCTU	4330

Table 40

2663	AUCCCAAU G UUAUAAA	1651	UUUAGUAA GCgaaagGCGaGuCaaGGuCu AUUGGGAU	4331
2717	CAGAGUAU G UAGUUAAU	1652	AUUAAUA GCgaaagGCGaGuCaaGGuCu AUACUCUG	4332
2901	AUCUUUCU G UCCCAAU	1653	AUUGGGA GCgaaagGCGaGuCaaGGuCu AGAAAGAU	4333
3071	GGGGACU G UUGGGUG	1654	CACCCAA GCgaaagGCGaGuCaaGGuCu AGUCCCC	4334
3111	UCACAAU G UGCCAGA	1655	UGCUGCA GCgaaagGCGaGuCaaGGuCu AGUUGUA	4335
40	AUCCAGA G UCAGGCC	1656	GGCCUGA GCgaaagGCGaGuCaaGGuCu UCUGGGAU	4336
46	GAGUCAG G CCCUGUAC	1657	GUACAGG GCgaaagGCGaGuCaaGGuCu CCUGACUC	4337
65	UCCUGUG G UGGUCCA	1658	UGGAGCA GCgaaagGCGaGuCaaGGuCu CAGCAGGA	4338
68	UGCUGUG G CUCCAGU	1659	AACUGAG GCgaaagGCGaGuCaaGGuCu CACCAGCA	4339
74	UGGCUCCA G UUCAGAA	1660	UCCUAAA GCgaaagGCGaGuCaaGGuCu UGGAGCCA	4340
85	CAGGAACA G UGAGCCU	1661	AGGCUCA GCgaaagGCGaGuCaaGGuCu UGUUCCUG	4341
89	AACAGUA G CCCUGUC	1662	GAGCAGG GCgaaagGCGaGuCaaGGuCu UCACUGUU	4342
120	GCUAUUC G UCAUUCU	1663	AAGAUUA GCgaaagGCGaGuCaaGGuCu GAUAUGGC	4343
196	CCUUGUC G UGUUACG	1664	CUGUACA GCgaaagGCGaGuCaaGGuCu GAGCAGG	4344
205	UGUACAG G CGGGUUU	1665	AAACCCG GCgaaagGCGaGuCaaGGuCu CUGUAACA	4345
210	CAGCGGG G UUUUUCU	1666	AAGAAAA GCgaaagGCGaGuCaaGGuCu CCCGCCUG	4346
248	ACCACAGA G UCUAGACU	1667	AGUCUAGA GCgaaagGCGaGuCaaGGuCu UCUGUGGU	4347
258	CUAGACUC G UGUUGAC	1668	GUCCACA GCgaaagGCGaGuCaaGGuCu GAGUCUAG	4348
261	GACUCUG G UGGACUC	1669	GAUGUCA GCgaaagGCGaGuCaaGGuCu CACGAGUC	4349
295	GAACACCC G UGUUGUCU	1670	AAGACACA GCgaaagGCGaGuCaaGGuCu GGGUGUUC	4350
305	GUGUCUG G CCAAAAU	1671	AAUUTUG GCgaaagGCGaGuCaaGGuCu CAAGACAC	4351
318	AAUUCGA G UCCCAAU	1672	AUUUGGA GCgaaagGCGaGuCaaGGuCu UGCGAAU	4352
332	AAUGUCA G UCACUCAC	1673	GUGAGUA GCgaaagGCGaGuCaaGGuCu UGGAGAU	4353
368	UUGUCCUG G UUAUCGU	1674	AGCGAUA GCgaaagGCGaGuCaaGGuCu CAGGACAA	4354
390	UGUUGCG G CGUUUAU	1675	AUAAAACG GCgaaagGCGaGuCaaGGuCu CGCAGACA	4355
392	UCUGCGC G UUUUAUA	1676	UGAUAUA GCgaaagGCGaGuCaaGGuCu GCCGCAGA	4356
442	UCUUGUG G UUCUUCU	1677	CAGAAGA GCgaaagGCGaGuCaaGGuCu CAACAAGA	4357
461	CUAUAAG G UAUGUUG	1678	GCAACAU GCgaaagGCGaGuCaaGGuCu CUUGAUAG	4358
472	UGUUGCC G UUUUCCU	1679	AGGACAA GCgaaagGCGaGuCaaGGuCu GGGCAACA	4359
506	AAACACA G CACCGAC	1680	GUCCGUG GCgaaagGCGaGuCaaGGuCu UGGUUGUU	4360
625	CAUCUUG G CUUUGCA	1681	UGCGAAG GCgaaagGCGaGuCaaGGuCu CCAAGAUG	4361
648	CUAUGGA G UGGGCCU	1682	GAGGCCA GCgaaagGCGaGuCaaGGuCu UCCCAUAG	4362
652	GGAGUGG G CCUCAGUC	1683	GACUGAG GCgaaagGCGaGuCaaGGuCu CCACUCCC	4363
658	GGGCCUA G UCCGUUUC	1684	GAAACGA GCgaaagGCGaGuCaaGGuCu UGAGGCC	4364

Table 40

662	CUCAGUCC	G	UUUCUCUU	1685	AAGAGAAA	GCCgaaagGCGaGuCaaGGuCu	GGACUGAG	4365
672	UUCUCUUG	G	CUCAGUUU	1686	AAACUGAG	GCCgaaagGCGaGuCaaGGuCu	CAAGAGAA	4366
677	UUGGCUCA	G	UUUACUAG	1687	CUAGUAAA	GCCgaaagGCGaGuCaaGGuCu	UGAGCCAA	4367
685	GUUUAUUA	G	UGCCAUTU	1688	AAAUUGCA	GCCgaaagGCGaGuCaaGGuCu	UAGUAAAC	4368
699	UUUGUUUA	G	UGGUUCGU	1689	ACGAACCA	GCCgaaagGCGaGuCaaGGuCu	UGAACAAA	4369
702	GUUCAGUG	G	UUCGUAGG	1690	CCUACGAA	GCCgaaagGCGaGuCaaGGuCu	CACUGAAC	4370
706	AGUGGUUC	G	UAGGGCUU	1691	AAGCCCUA	GCCgaaagGCGaGuCaaGGuCu	GAACCCAU	4371
711	UUCGUAGG	G	CUUUCGCC	1692	GGGAAAG	GCCgaaagGCGaGuCaaGGuCu	CCUACGAA	4372
729	ACTUGUCU	G	CUUUCAGU	1693	ACUGAAAG	GCCgaaagGCGaGuCaaGGuCu	CAGACAGU	4373
736	GGCUUUUA	G	UUUAUUGG	1694	CCAUAUAA	GCCgaaagGCGaGuCaaGGuCu	UGAAAGCC	4374
753	AUGAUGUG	G	UUUUGGGG	1695	CCCCAAAA	GCCgaaagGCGaGuCaaGGuCu	CACAUAU	4375
762	UUUUGGGG	G	CCAAGUCU	1696	AGACUUUG	GCCgaaagGCGaGuCaaGGuCu	CCCCAAAA	4376
767	GGGCCCCA	G	UCUGUACA	1697	UGUACAGA	GCCgaaagGCGaGuCaaGGuCu	UUGGCCCC	4377
785	CAUCUUGA	G	UCCUUUUA	1698	UAAAGGGA	GCCgaaagGCGaGuCaaGGuCu	UCAAGAUG	4378
826	GUCUUUGG	G	UAUACAUU	1699	AAUGUAUA	GCCgaaagGCGaGuCaaGGuCu	CCAAAGAC	4379
898	AAUUGGGA	G	UUGGGGCA	1700	UGCCCCAA	GCCgaaagGCGaGuCaaGGuCu	UCCCAAUU	4380
904	GAGUUGGG	G	CACAUUGC	1701	GCAUUGUG	GCCgaaagGCGaGuCaaGGuCu	CCCAACUC	4381
971	GUAAACAG	G	CCAUUUGA	1702	UCAAUAGG	GCCgaaagGCGaGuCaaGGuCu	CUGUUUAC	4382
987	AUUGGAAA	G	UAUGUCAA	1703	UUGACAUU	GCCgaaagGCGaGuCaaGGuCu	UUUCCAAU	4383
1006	AAUUGUGG	G	UCUUUUGG	1704	CCAAAGA	GCCgaaagGCGaGuCaaGGuCu	CCCAAUU	4384
1016	UUUUUGGG	G	UUUGCCGC	1705	GCGGCAAA	GCCgaaagGCGaGuCaaGGuCu	CCCCAAAG	4385
1080	GCAUACAA	G	CAAAACAG	1706	CUGUUUUG	GCCgaaagGCGaGuCaaGGuCu	UUGUAUGC	4386
1089	CAAAACAG	G	CUUUUACU	1707	AGUAAAAG	GCCgaaagGCGaGuCaaGGuCu	CUGUUUUG	4387
1116	CUUACAAG	G	CCUUUCUA	1708	UAGAAAGG	GCCgaaagGCGaGuCaaGGuCu	CUUGUAAG	4388
1126	CUUUCUAA	G	UAAACAGU	1709	ACUGUUUA	GCCgaaagGCGaGuCaaGGuCu	UUAGAAAG	4389
1133	AGUAAACA	G	UAUGUGAA	1710	UUCACAUU	GCCgaaagGCGaGuCaaGGuCu	UGUUUACU	4390
1152	UUUACCCC	G	UUGUCUGG	1711	CCGAGCAA	GCCgaaagGCGaGuCaaGGuCu	GGGUUAAA	4391
1160	GUUGCUCG	G	CAACGGCC	1712	GGCCUUG	GCCgaaagGCGaGuCaaGGuCu	CGAGCAAC	4392
1166	CGGCAACG	G	CCUGGUUCU	1713	AGACCAGG	GCCgaaagGCGaGuCaaGGuCu	CGUUGCCG	4393
1171	ACGGCCUG	G	UCUUGGCC	1714	GGCAUAGA	GCCgaaagGCGaGuCaaGGuCu	CAGGCCGU	4394
1182	UAUGCCAA	G	UGUUGUCU	1715	AGCAAAAC	GCCgaaagGCGaGuCaaGGuCu	UUGGCAUA	4395
1207	CCCAACUG	G	UUGGGGCU	1716	AGCCCAA	GCCgaaagGCGaGuCaaGGuCu	CAGUGGGG	4396
1213	UGGUUGGG	G	CUUGGCCA	1717	UGGCCAAG	GCCgaaagGCGaGuCaaGGuCu	CCCAACCA	4397
1218	GGGCUUG	G	CCAUAGGC	1718	GCCUAUGG	GCCgaaagGCGaGuCaaGGuCu	CAAGCCCC	4398

Table 40

1225	GGCCAUG	G	CCAUCAGC	1719	GCUGAUGG	GCgaaagGCGaGuCaaGGuCu	CUAUGGCC	4399
1232	GGCCAUC	A	GCGAUGG	1720	CGCAUGCG	GCgaaagGCGaGuCaaGGuCu	UGAUGGCC	4400
1240	GCGCAUG	C	UGGAACCU	1721	AGGUUCCA	GCgaaagGCGaGuCaaGGuCu	GCAUGCGC	4401
1287	AACUCCU	A	GCGUUGU	1722	ACAAGCGG	GCgaaagGCGaGuCaaGGuCu	UAGGAGUU	4402
1306	UGCUGCA	G	CAGGUUG	1723	CAGACCUG	GCgaaagGCGaGuCaaGGuCu	UGCAGACA	4403
1310	CGCAGCA	G	UCUGGGC	1724	GCCCCAGA	GCgaaagGCGaGuCaaGGuCu	CUGCUGCG	4404
1317	GGUCUGG	G	CAAAACUC	1725	GAGUUJUG	GCgaaagGCGaGuCaaGGuCu	CCCAGACC	4405
1347	AUUCUGU	C	UGCUCUC	1726	GGAGAGCA	GCgaaagGCGaGuCaaGGuCu	GACAGAAU	4406
1379	UUUCCAUG	G	CUGCUAGG	1727	CCUAGCAG	GCgaaagGCGaGuCaaGGuCu	CAUGGAAA	4407
1387	GCUGCUAG	G	CUGUGCUG	1728	CAGCACAG	GCgaaagGCGaGuCaaGGuCu	CUAGCAGC	4408
1418	CGCGGAC	G	UCCUUUGU	1729	ACAAAGGA	GCgaaagGCGaGuCaaGGuCu	GUCCCGCG	4409
1431	UUGUUUAC	G	UCCCGUCG	1730	CGACGGGA	GCgaaagGCGaGuCaaGGuCu	GUAAACAA	4410
1436	UACGUCCC	G	UCGGCGCU	1731	AGCGCCGA	GCgaaagGCGaGuCaaGGuCu	GGGACGUA	4411
1440	UCCGUCG	G	CGCUGAAU	1732	AUUCAGCG	GCgaaagGCGaGuCaaGGuCu	CGACGGGA	4412
1471	CUCCCGG	G	CCGUUGG	1733	CCAAGCGG	GCgaaagGCGaGuCaaGGuCu	CCCAGGAG	4413
1481	CGCUUGG	G	CUCUACCG	1734	CGGUAGAG	GCgaaagGCGaGuCaaGGuCu	CCCAGCGG	4414
1517	UACCGACC	G	UCCACGGG	1735	CCCUGUGA	GCgaaagGCGaGuCaaGGuCu	GGUCGGUA	4415
1526	UCCACGGG	G	CGACCTUC	1736	GAGUGUGG	GCgaaagGCGaGuCaaGGuCu	CCCUGUGA	4416
1553	GACUCCCC	G	UCUGUGCC	1737	GGCACAGA	GCgaaagGCGaGuCaaGGuCu	GGGAGUUC	4417
1579	GCCGACC	G	UGUGCACU	1738	AGUGCACA	GCgaaagGCGaGuCaaGGuCu	GGUCCGGC	4418
1605	CUCUGCAC	G	UCGCAUGG	1739	CCAUGCGA	GCgaaagGCGaGuCaaGGuCu	GUGCAGAG	4419
1622	AGACCACC	G	UGAACGCC	1740	GGCGUUCA	GCgaaagGCGaGuCaaGGuCu	GGUGGUCU	4420
1649	UGCCCAAG	G	UCUUGCAU	1741	AUGCAAGA	GCgaaagGCGaGuCaaGGuCu	CUUGGGCA	4421
1679	GACUUUCA	G	CAUUGUCA	1742	UGACAUUG	GCgaaagGCGaGuCaaGGuCu	UGAAAGUC	4422
1703	ACCUUGAG	G	CAUACUUC	1743	GAAGUAUG	GCgaaagGCGaGuCaaGGuCu	CUCAGGUA	4423
1732	UUUAAUGA	G	UGGGAGGA	1744	UCCUCCCA	GCgaaagGCGaGuCaaGGuCu	UCAUUAAA	4424
1741	UGGAGGGA	G	UUGGGGGA	1745	UCCCCCAA	GCgaaagGCGaGuCaaGGuCu	UCCUCCCA	4425
1754	GGGAGGAG	G	UUAGGUUA	1746	UAACCUAA	GCgaaagGCGaGuCaaGGuCu	CUCCUCCC	4426
1759	GAGGUUAG	G	UUAAAGGU	1747	ACCUUUA	GCgaaagGCGaGuCaaGGuCu	CUAACCTUC	4427
1766	GGUUAAG	G	UCUUUGUA	1748	UACAAAGA	GCgaaagGCGaGuCaaGGuCu	CUUUAACC	4428
1782	ACUAGGAG	G	CUGUAGGC	1749	GCCUACAG	GCgaaagGCGaGuCaaGGuCu	CUCCUAGU	4429
1789	GGCUGUAG	G	CAUAAAUU	1750	AUUUAUG	GCgaaagGCGaGuCaaGGuCu	CUACAGCC	4430
1799	AUAAAUUG	G	UGUGUUCA	1751	UGAACACA	GCgaaagGCGaGuCaaGGuCu	CAUUUAU	4431
1811	GUUCACCA	G	CACCAUGC	1752	GCAUGGUG	GCgaaagGCGaGuCaaGGuCu	UGGUGAAC	4432

Table 40

1870	CUGUUCAA	G	CCUCCAAAG	1753	CUUGGAGG	GCCgaaagGCCgGaGuCaaGGGuCu	UUGAAACAG	4433
1878	GCCUCCAA	G	CUGUGCCU	1754	AGGCACAG	GCCgaaagGCCgGaGuCaaGGGuCu	UUGGAGGC	4434
1890	UGCCUUGG	G	UGGCUUUG	1755	CAAAAGCCA	GCCgaaagGCCgGaGuCaaGGGuCu	CCAAGGCA	4435
1893	CUTUGGUG	G	CUUUGGGG	1756	CCCCAAAG	GCCgaaagGCCgGaGuCaaGGGuCu	CACCCAAAG	4436
1901	GCUTUGGG	G	CAUGGACA	1757	UGUCCAU	GCCgaaagGCCgGaGuCaaGGGuCu	CCCAAAAGC	4437
1917	AUTGACCC	G	UAUAAAGA	1758	UCUUAUA	GCCgaaagGCCgGaGuCaaGGGuCu	GGGUCAAU	4438
1933	AAUTUGGA	G	CUUCUGUG	1759	CACAGAAG	GCCgaaagGCCgGaGuCaaGGGuCu	UCCAAAUU	4439
1944	UCUGUGGA	G	UUACUCUC	1760	GAGAGUAA	GCCgaaagGCCgGaGuCaaGGGuCu	UCCACAGA	4440
2023	AUCGGGGG	G	CCUAGAG	1761	CUCUAAGG	GCCgaaagGCCgGaGuCaaGGGuCu	CCCCCGAU	4441
2031	GCCUAGA	G	UCUCCGGA	1762	UCCGGAGA	GCCgaaagGCCgGaGuCaaGGGuCu	UCUAAGGC	4442
2062	ACCAUACG	G	CACUCAGG	1763	CCUGAGUG	GCCgaaagGCCgGaGuCaaGGGuCu	CGUAUGGU	4443
2070	GCACUCAG	G	CAAGCUAU	1764	AUAGCUUG	GCCgaaagGCCgGaGuCaaGGGuCu	CUGAGUGC	4444
2074	UCAGGCAA	G	CUAUUCUG	1765	CAGAAUAG	GCCgaaagGCCgGaGuCaaGGGuCu	UUGCCUGA	4445
2090	GUGUUGGG	G	UGAUUGA	1766	UCAACUCA	GCCgaaagGCCgGaGuCaaGGGuCu	CCCAACAC	4446
2094	UGGGGUGA	G	UUGAUGAA	1767	UUAUCUA	GCCgaaagGCCgGaGuCaaGGGuCu	UCACCCCA	4447
2107	UGAAUCUA	G	CCACCUGG	1768	CCAGUGG	GCCgaaagGCCgGaGuCaaGGGuCu	UAGAUAUA	4448
2116	CCACCUGG	G	UGGGAAGU	1769	ACUCCCCA	GCCgaaagGCCgGaGuCaaGGGuCu	CCAGGUGG	4449
2123	GGUGGGAA	G	UAAUUUGG	1770	CCAAUAUA	GCCgaaagGCCgGaGuCaaGGGuCu	UUCCCACC	4450
2140	AAGAUCCA	G	CAUCCAGG	1771	CCUGGAG	GCCgaaagGCCgGaGuCaaGGGuCu	UGGAUCUU	4451
2155	GGGAUAUA	G	UAGUCAGC	1772	GCUGACTUA	GCCgaaagGCCgGaGuCaaGGGuCu	UAAUUCUU	4452
2158	AAUAUAUA	G	UCAGCUAU	1773	AUAGCUUA	GCCgaaagGCCgGaGuCaaGGGuCu	UACUAUAU	4453
2162	AGUAGUCA	G	CUAUGUCA	1774	UGACAUAG	GCCgaaagGCCgGaGuCaaGGGuCu	UGACUACU	4454
2173	AUGUCAAC	G	UUUAUAUG	1775	CAUAUAUA	GCCgaaagGCCgGaGuCaaGGGuCu	GUUGACAU	4455
2183	UAAUAUGG	G	CCUAAAAA	1776	UUUUUAGG	GCCgaaagGCCgGaGuCaaGGGuCu	CCAUAUAUA	4456
2208	CUAUUUG	G	UUUCACAU	1777	AUGUGAAA	GCCgaaagGCCgGaGuCaaGGGuCu	CACAAUAG	4457
2235	ACUUUUGG	G	CGAGAAAC	1778	GUUUCUG	GCCgaaagGCCgGaGuCaaGGGuCu	CCAAAAGU	4458
2260	AAUAUUUG	G	UGUCUUUU	1779	AAAAGACA	GCCgaaagGCCgGaGuCaaGGGuCu	CRAUAUAU	4459
2272	CUUUUGGA	G	UGUGGAUU	1780	AAUCCACA	GCCgaaagGCCgGaGuCaaGGGuCu	UCCAAAAG	4460
2360	ACGAAGAG	G	CAGGUCCC	1781	GGGACCU	GCCgaaagGCCgGaGuCaaGGGuCu	CUCUUUGU	4461
2364	AGAGGCAG	G	UCCCUUAG	1782	CUAGGGGA	GCCgaaagGCCgGaGuCaaGGGuCu	CUGCCUCU	4462
2403	AGACGAAG	G	UCUCAUUC	1783	GAUUGAGA	GCCgaaagGCCgGaGuCaaGGGuCu	CUUCGUCU	4463
2417	AUCGCCGC	G	UCGCAGAA	1784	UUCUGCGA	GCCgaaagGCCgGaGuCaaGGGuCu	GCAGCGAU	4464
2454	CRAUGUUA	G	UAUUCUUU	1785	AAGGAAUA	GCCgaaagGCCgGaGuCaaGGGuCu	UAACAUUG	4465
2474	CACAUAAAG	G	UGGGAAC	1786	GUUUCUCA	GCCgaaagGCCgGaGuCaaGGGuCu	CUUAUGUG	4466

Table 40

2491	UUUACGGG G CUUUAUUC	1787	GAUUAAG GCgaaagGCGaGuCaaGGuCu	CCC GUAAA	4467
2507	CUUCUACG G UACCUUGC	1788	GCAAGGUA GCgaaagGCGaGuCaaGGuCu	CGUAGAAG	4468
2530	CCUAAUG G CAAACUCC	1789	GGAGUTUG GCgaaagGCGaGuCaaGGuCu	CAUJUAGG	4469
2587	AGAUGUAA G CAAUUUGU	1790	ACAAUUTG GCgaaagGCGaGuCaaGGuCu	UUACAUCU	4470
2599	UUUGUGGG G CCCCUUAC	1791	GUAGGGG GCgaaagGCGaGuCaaGGuCu	CCCACAAA	4471
2609	CCCUUACA G UAAUUGAA	1792	UUCAUUA GCgaaagGCGaGuCaaGGuCu	UGUAAGGG	4472
2650	CCUGCUAG G UUUUAUCC	1793	GGAUAAA GCgaaagGCGaGuCaaGGuCu	CUAGCAGG	4473
2701	AUCAACC G UAUUAUCC	1794	GGAUAAU GCgaaagGCGaGuCaaGGuCu	GGUUUGAU	4474
2713	UAUCCAGA G UAUUGAGU	1795	ACUACAU GCgaaagGCGaGuCaaGGuCu	UCUGGAUA	4475
2720	AGUAUGUA G UUAUUCAU	1796	AUGAUUA GCgaaagGCGaGuCaaGGuCu	UACAUACU	4476
2768	UUUGGAAG G CGGGGAUC	1797	GAUCCCG GCgaaagGCGaGuCaaGGuCu	CUUCCAAA	4477
2791	AAAAGAGA G UCCACACG	1798	CGUGUGGA GCgaaagGCGaGuCaaGGuCu	UCUCUUUU	4478
2799	GUCCACAC G UAGCGCCU	1799	AGGCGCUA GCgaaagGCGaGuCaaGGuCu	GUGUGGAC	4479
2802	CACACGUA G CGCCUCAU	1800	AUGAGCG GCgaaagGCGaGuCaaGGuCu	UACGUGUG	4480
2818	UUUGCGG G UCACCAUA	1801	UAUGUGA GCgaaagGCGaGuCaaGGuCu	CCGCAAAA	4481
2848	GAUCUACA G CAUGGGAG	1802	CUCCCAUG GCgaaagGCGaGuCaaGGuCu	UGUAGAUC	4482
2857	CAUGGAG G UUGGUCUU	1803	AAGACAA GCgaaagGCGaGuCaaGGuCu	CUCCCAUG	4483
2861	GGAGUUG G UCUUCCAA	1804	UUGGAAGA GCgaaagGCGaGuCaaGGuCu	CAACCUCC	4484
2881	UCGAAAAG G CAUGGGGA	1805	UCCCAUG GCgaaagGCGaGuCaaGGuCu	CUUUUGGA	4485
2936	GAUCAUA G UUGGACCC	1806	GGGUCAA GCgaaagGCGaGuCaaGGuCu	UGAUGAUC	4486
2955	CAUCAA G CCAACUCA	1807	UGAGUUG GCgaaagGCGaGuCaaGGuCu	UUUGAUG	4487
2964	CCAACUA G UAAAUCCA	1808	UGAUUUA GCgaaagGCGaGuCaaGGuCu	UGAGUUGG	4488
3005	GACACUG G CCGGACGC	1809	CGUCCGG GCgaaagGCGaGuCaaGGuCu	CAGUUGUC	4489
3021	CCAACAAG G UGGGAGUG	1810	CACUCCA GCgaaagGCGaGuCaaGGuCu	CUUUGUGG	4490
3027	AGGUGGA G UGGGAGCA	1811	UGUCCCA GCgaaagGCGaGuCaaGGuCu	UCCCACTU	4491
3033	GAGUGGA G CAUUCGGG	1812	CCCGAUG GCgaaagGCGaGuCaaGGuCu	UCCCACTU	4492
3041	GCAUUCG G CCAGGGUU	1813	AACCUUG GCgaaagGCGaGuCaaGGuCu	CCGAAUGC	4493
3047	GGGCCAGG G UUCACCCC	1814	GGGUGAA GCgaaagGCGaGuCaaGGuCu	CCUGGCCC	4494
3077	CUGUUGG G UGGAGCCC	1815	GGGUCCA GCgaaagGCGaGuCaaGGuCu	CCCAACAG	4495
3082	GGGUGGA G CCCUCACG	1816	CGUGAGG GCgaaagGCGaGuCaaGGuCu	UCCACCCC	4496
3097	CGCUCAGG G CCUACUCA	1817	UGAGUAG GCgaaagGCGaGuCaaGGuCu	CCUGAGCG	4497
3117	CUGUGCCA G CAGUCCU	1818	AGGAGUG GCgaaagGCGaGuCaaGGuCu	UGGCACAG	4498
3120	UGCCAGCA G CUCCUCCU	1819	AGGAGAG GCgaaagGCGaGuCaaGGuCu	UGCUGGCA	4499
3146	ACCAAUCG G CAGUCAGG	1820	CCUGACUG GCgaaagGCGaGuCaaGGuCu	CGAUUGGU	4500

Table 40

3149	AAUCGGCA G UCAGGAAG	1821	CUUCCUGA GCCgaaagGCCaGuCaagGuCu	UGCCGAUU	4501
3158	UCAGGAAG G CAGCCUAC	1822	GUAGGCUG GCCgaaagGCCaGuCaagGuCu	CUUCCUGA	4502
3161	GGAAGGCA G CCUACUCC	1823	GGAGUAGG GCCgaaagGCCaGuCaagGuCu	UGCCUJCC	4503
3204	AUCCUCAG G CCAUGCAG	1824	CUGCAUGG GCCgaaagGCCaGuCaagGuCu	CUGAGGAU	4504

Input Sequence = AF100308. Cut Site = YG/M or UG/U.
Stem Length = 8 . Core Sequence = GCCgaaagGCCaGuCaagGuCu
AF100308 (Hepatitis B virus strain 2-18, 3215 bp)

Table 41

Table 41: Human HBV DNzyme and Substrate Sequence

Pos	Substrate	Seq ID	DNzyme	Rx Seq ID
508	CAACCAGC A CCGGACCA	833	TGGTCCGG GGCTAGCTACAACGA GCTGGTTG	4505
1632	GAACGCCC A CAGGAACC	1096	GGTTCCTG GGCTAGCTACAACGA GGGCGTTC	4506
2992	CAACCCGC A CAAGGACA	1376	TGTCCTTG GGCTAGCTACAACGA GCGGGTTG	4507
61	ACUUUCCU G CUGGUGGC	1448	GCCACCAG GGCTAGCTACAACGA AGGAAAGT	4508
94	UGAGCCCU G CUCAGAAU	1450	ATTCTGAG GGCTAGCTACAACGA AGGGCTCA	4509
112	CUGUCUCU G CCAUAUCG	1451	CGATATGG GGCTAGCTACAACGA AGAGACAG	4510
169	AGAACAUC G CAUCAGGA	1454	TCCTGATG GGCTAGCTACAACGA GATGTTCT	4511
192	GGACCCCU G CUCGUGUU	1455	AACACGAG GGCTAGCTACAACGA AGGGGTCC	4512
315	CAAAAUUC G CAGUCCCA	1457	TGGGACTG GGCTAGCTACAACGA GAATTTTG	4513
374	UGGUUAUC G CUGGAUGU	1458	ACATCCAG GGCTAGCTACAACGA GATAACCA	4514
387	AUGUGUCU G CGGCGUUU	1459	AAACGCCG GGCTAGCTACAACGA AGACACAT	4515
410	CUUCCUCU G CAUCCUGC	1460	GCAGGATG GGCTAGCTACAACGA AGAGGAAG	4516
417	UGCAUCCU G CUGCUAUG	1461	CATAGCAG GGCTAGCTACAACGA AGGATGCA	4517
420	AUCCUGCU G CUAUGCCU	1462	AGGCATAG GGCTAGCTACAACGA AGCAGGAT	4518
425	GCUGCUAU G CCUCAUCU	1463	AGATGAGG GGCTAGCTACAACGA ATAGCAGC	4519
468	GGUAUGUU G CCCGUUUG	1464	CAAACGGG GGCTAGCTACAACGA AACATACC	4520
518	CGGACCAU G CAAAACCU	1465	AGGTTTTG GGCTAGCTACAACGA ATGGTCCG	4521
527	CAAAACCU G CACAACUC	1466	GAGTTGTG GGCTAGCTACAACGA AGGTTTTG	4522
538	CAACUCCU G CUCAAGGA	1467	TCCTTGAG GGCTAGCTACAACGA AGGAGTTG	4523
569	CUCAUGUU G CUGUACAA	1468	TTGTACAG GGCTAGCTACAACGA AACATGAG	4524
596	CGGAAACU G CACCUGUA	1469	TACAGGTG GGCTAGCTACAACGA AGTTTCCG	4525
631	GGGCUUUC G CAAAUAC	1470	GTATTTTG GGCTAGCTACAACGA GAAAGCCC	4526
687	UUACUAGU G CCAUUUGU	1471	ACAAATGG GGCTAGCTACAACGA ACTAGTAA	4527
795	CCUUUUAU G CCGCUGUU	1474	AACAGCGG GGCTAGCTACAACGA ATAAAGGG	4528
798	UUUAUGCC G CUGUUACC	1475	GGTAACAG GGCTAGCTACAACGA GGCATAAA	4529
911	GGCAUAU G CCACAGGA	1476	TCCTGTGG GGCTAGCTACAACGA AATGTGCC	4530
1020	UGGGGUUU G CGGCCCUU	1479	AGGGGCGG GGCTAGCTACAACGA AAACCCCA	4531
1023	GGUUUGCC G CCCCUUUC	1480	GAAAGGGG GGCTAGCTACAACGA GGCAAACC	4532
1034	CCUUUCAC G CAUUGUGG	1481	CCACATTG GGCTAGCTACAACGA GTGAAAGG	4533
1050	GAUAUUCU G CUUUAUUG	1482	CATTAAAG GGCTAGCTACAACGA AGAATATC	4534
1058	GCUUUAU G CCUUUAUA	1483	TATAAAGG GGCTAGCTACAACGA ATTAAAGC	4535
1068	CUUUAUUA G CAUGCAUA	1484	TATGCATG GGCTAGCTACAACGA ATATAAAG	4536
1072	AUAUGCAU G CAUACAAG	1485	CTTGTATG GGCTAGCTACAACGA ATGCATAT	4537
1103	ACUUUCUC G CCAACUUA	1486	TAAGTTGG GGCTAGCTACAACGA GAGAAAGT	4538
1155	ACCCCGUU G CUCGGCAA	1488	TTGCCGAG GGCTAGCTACAACGA AACGGGGT	4539
1177	UGGUCUAU G CCAAGUGU	1489	ACACTTGG GGCTAGCTACAACGA ATAGACCA	4540
1188	AAGUGUUU G CUGACGCA	1490	TGCGTCAG GGCTAGCTACAACGA AAACACTT	4541
1194	UUGCUGAC G CAACCCCC	1492	GGGGGTTG GGCTAGCTACAACGA GTCAGCAA	4542
1234	CCAUCAGC G CAUGCGUG	1493	CACGCATG GGCTAGCTACAACGA GCTGATGG	4543
1238	CAGCGCAU G CGUGGAAC	1494	GTTCCACG GGCTAGCTACAACGA ATGCGCTG	4544
1262	UCUCCUCU G CCGAUCCA	1495	TGGATCGG GGCTAGCTACAACGA AGAGGAGA	4545
1275	UCCAUAAC G CGGAACUC	1497	GAGTTCCG GGCTAGCTACAACGA GGTATGGA	4546
1290	UCCUAGCC G CUUGUUUU	1498	AAAACAAG GGCTAGCTACAACGA GGCTAGGA	4547
1299	CUUGUUUU G CUCGCAGC	1499	GCTGCGAG GGCTAGCTACAACGA AAAACAAG	4548
1303	UUUUGCUC G CAGCAGGU	1500	ACCTGCTG GGCTAGCTACAACGA GAGCAAAA	4549
1349	UCUGUCGU G CUCUCCCG	1502	CGGGAGAG GGCTAGCTACAACGA ACGACAGA	4550
1357	GCUCUCCC G CAAUAUAU	1503	TATATTTG GGCTAGCTACAACGA GGGAGAGC	4551

Table 41

1382	CCAUGGCU G CUAGGCU	1504	CAGCCTAG GGCTAGCTACAACGA AGCCATGG	4552
1392	UAGGCGUG G CUGCCAAC	1505	GTTGGCAG GGCTAGCTACAACGA ACAGCCTA	4553
1395	GCUGUGCU G CCAACUGG	1506	CCAGTTGG GGCTAGCTACAACGA AGCACAGC	4554
1411	GAUCCUAC G CGGGACGU	1507	ACGTCCCG GGCTAGCTACAACGA GTAGGATC	4555
1442	CCGUCGGC G CUGAAUCC	1508	GGATTCAG GGCTAGCTACAACGA GCCGACGG	4556
1452	UGAAUCCC G CGGACGAC	1510	GTCGTCCG GGCTAGCTACAACGA GGGATTCA	4557
1474	CCGGGGCC G CUUGGGGC	1512	GCCCAAG GGCTAGCTACAACGA GGCCCCGG	4558
1489	GCUCUACC G CCCGCUUC	1513	GAAGCGGG GGCTAGCTACAACGA GGTAGAGC	4559
1493	UACCGCCC G CUUCUCCG	1514	CGGAGAAG GGCTAGCTACAACGA GGGCGGTA	4560
1501	GCUCUCC G CCUAUUGU	1515	ACAATAGG GGCTAGCTACAACGA GGAGAAGC	4561
1528	CACGGGGC G CACCUCUC	1517	GAGAGGTG GGCTAGCTACAACGA GCCCCGTG	4562
1542	CUCUUUAC G CGGACUCC	1518	GGAGTCCG GGCTAGCTACAACGA GTAAAGAG	4563
1559	CCGUCUGU G CCUUCUCA	1519	TGAGAAGG GGCTAGCTACAACGA ACAGACGG	4564
1571	UCUCAUCU G CCGGACCG	1520	CGGTCCGG GGCTAGCTACAACGA AGATGAGA	4565
1583	GACCGUGU G CACUUCGC	1521	GCGAAGTG GGCTAGCTACAACGA ACACGGTC	4566
1590	UGCACUUC G CUUCACCU	1522	AGGTGAAG GGCTAGCTACAACGA GAAGTGCA	4567
1601	UCACCUCU G CACGUCGC	1523	GCGACGTG GGCTAGCTACAACGA AGAGGTGA	4568
1608	UGCACGUC G CAUGGAGA	1524	TCTCCATG GGCTAGCTACAACGA GACGTGCA	4569
1628	CCGUGAAC G CCCACAGG	1526	CCTGTGGG GGCTAGCTACAACGA GTTCACGG	4570
1642	AGGAACCU G CCCAAGGU	1527	ACCTTGGG GGCTAGCTACAACGA AGGTTCTT	4571
1654	AAGGUCUU G CAUAAGAG	1528	CTCTTATG GGCTAGCTACAACGA AAGACCTT	4572
1818	AGCACCAU G CAACUUUU	1533	AAAAGTTG GGCTAGCTACAACGA ATGGTGCT	4573
1835	UCACCUCU G CCUAUUA	1534	TGATTAGG GGCTAGCTACAACGA AGAGGTGA	4574
1883	CAAGCUGU G CCUUGGGU	1535	ACCCAAGG GGCTAGCTACAACGA ACAGCTTG	4575
1959	UCUUUUUU G CCUUCUGA	1537	TCAGAAGG GGCTAGCTACAACGA AAAAAAGA	4576
2002	UCGACACC G CCUCUGCU	1541	AGCAGAGG GGCTAGCTACAACGA GGTGTCGA	4577
2008	CCGCCUCU G CUCUGUAU	1542	ATACAGAG GGCTAGCTACAACGA AGAGGCGG	4578
2282	GUGGAUUC G CACUCCUC	1548	GAGGAGTG GGCTAGCTACAACGA GAATCCAC	4579
2293	CUCCUCCU G CAUAUAGA	1549	TCTATATG GGCTAGCTACAACGA AGGAGGAG	4580
2311	CACCAAU G CCCUAUC	1550	GATAGGGG GGCTAGCTACAACGA ATTTGGTG	4581
2388	ACUCCUC G CCUCGCAG	1552	CTGCGAGG GGCTAGCTACAACGA GAGGGAGT	4582
2393	CUCGCCUC G CAGACGAA	1553	TTCGTCTG GGCTAGCTACAACGA GAGGCGAG	4583
2412	UCUCAUUC G CCGCGUCG	1555	CGACGCGG GGCTAGCTACAACGA GATTGAGA	4584
2415	CAUUCGCC G CGUCGCAG	1556	CTGCGACG GGCTAGCTACAACGA GCGATTG	4585
2420	GCCGCGUC G CAGAAGAU	1557	ATCTTCTG GGCTAGCTACAACGA GACGCGGC	4586
2514	GGUACCUU G CUUUAUUC	1558	GATTAAAG GGCTAGCTACAACGA AAGGTACC	4587
2560	AUUCAUUU G CAGGAGGA	1560	TCCTCCTG GGCTAGCTACAACGA AAATGAAT	4588
2641	UUAACUUAU G CCUGCUAG	1563	CTAGCAGG GGCTAGCTACAACGA ATAGTTAA	4589
2645	CUAUGCCU G CUAGGUUU	1564	AAACCTAG GGCTAGCTACAACGA AGGCATAG	4590
2677	AAAUUUUU G CCCUAGA	1565	TCTAAGGG GGCTAGCTACAACGA AAATATTT	4591
2740	UCCAGAC G CGACAUUA	1566	TAATGTCTG GGCTAGCTACAACGA GTCTGGAA	4592
2804	CACGUAGC G CCUCAUUU	1568	AAATGAGG GGCTAGCTACAACGA GCTACGTG	4593
2814	CUCAUUUU G CGGGUCAC	1569	GTGACCCG GGCTAGCTACAACGA AAAATGAG	4594
2946	UGGACCCU G CAUUCAAA	1572	TTTGAATG GGCTAGCTACAACGA AGGGTCCA	4595
2990	CUCAACCC G CACAAGGA	1573	TCCTTGTG GGCTAGCTACAACGA GGGTTGAG	4596
3012	GGCCGGAC G CCAACAAG	1574	CTGTGTTG GGCTAGCTACAACGA GTCCGGCC	4597
3090	GCCCUCAC G CUCAGGGC	1575	GCCCTGAG GGCTAGCTACAACGA GTGAGGGC	4598
3113	ACAACUGU G CCAGCAGC	1576	GCTGCTGG GGCTAGCTACAACGA ACAGTTGT	4599
3132	CUCCUCCU G CCUCCACC	1577	GGTGGAGG GGCTAGCTACAACGA AGGAGGAG	4600
51	AGGGCCCU G UACUUUCC	1578	GGAAAGTA GGCTAGCTACAACGA AGGGCCCT	4601
106	AGAAUACU G UCUCUGCC	1579	GGCAGAGA GGCTAGCTACAACGA AGTATTCT	4602

Table 41

148	GGGACCCU G UACCGAAC	1580	GTTCGGTA GGCTAGCTACAACGA AGGGTCCC	4603
198	CUGCUCGU G UUACAGGC	1581	GCCTGTAA GGCTAGCTACAACGA ACGAGCAG	4604
219	UUUUUCUU G UUGACAAA	1582	TTTGTCAA GGCTAGCTACAACGA AAGAAAAA	4605
297	ACACCCGU G UGUCUUGG	1583	CCAAGACA GGCTAGCTACAACGA ACGGGTGT	4606
299	ACCCGUGU G UCUUGGCC	1584	GGCCAAGA GGCTAGCTACAACGA ACACGGGT	4607
347	ACCAACCU G UUGUCCUC	1585	GAGGACAA GGCTAGCTACAACGA AGGTTGGT	4608
350	AACCUGUU G UCCUCCAA	1586	TTGGAGGA GGCTAGCTACAACGA AACAGGTT	4609
362	UCCAAUUU G UCCUGGUU	1587	AACCAGGA GGCTAGCTACAACGA AAATTGGA	4610
381	CGCUGGAU G UGUCUGCG	1588	CGCAGACA GGCTAGCTACAACGA ATCCAGCG	4611
383	CUGGAUGU G UCUGCGGC	1589	GCCGAGA GGCTAGCTACAACGA ACATCCAG	4612
438	AUCUUCUU G UUGGUUCU	1590	AGAACCAA GGCTAGCTACAACGA AAGAAGAT	4613
465	CAAGGUAU G UUGCCCGU	1591	ACGGGCAA GGCTAGCTACAACGA ATACCTTG	4614
476	GCCCGUUU G UCCUCUAA	1592	TTAGAGGA GGCTAGCTACAACGA AAACGGGC	4615
555	ACCUCUAU G UUUCCUC	1593	GAGGGAAA GGCTAGCTACAACGA ATAGAGGT	4616
566	UCCCUCAU G UUGCUGUA	1594	TACAGCAA GGCTAGCTACAACGA ATGAGGGA	4617
572	AUGUUGCU G UACAAAC	1595	GTTTGTGA GGCTAGCTACAACGA AGCAACAT	4618
602	CUGCACCU G UAUUCCCA	1596	TGGGAATA GGCTAGCTACAACGA AGGTGCAG	4619
694	UGCCAUUU G UUCAGUGG	1597	CCACTGAA GGCTAGCTACAACGA AAATGGCA	4620
724	CCCCACU G UCUGGCUU	1598	AAGCCAGA GGCTAGCTACAACGA AGTGGGGG	4621
750	UGGAUGAU G UGGUUUUG	1599	CAAAACCA GGCTAGCTACAACGA ATCATCCA	4622
771	CCAAGUCU G UACAACAU	1600	ATGTTGTA GGCTAGCTACAACGA AGACTTGG	4623
801	AUGCCGCU G UUACCAU	1601	ATTGGTAA GGCTAGCTACAACGA AGCGGCAT	4624
818	UUUCUUUU G UCUUUGGG	1602	CCCAAAGA GGCTAGCTACAACGA AAAAGAAA	4625
888	UGGGAUUAU G UAAUUGGG	1603	CCCAATTA GGCTAGCTACAACGA ATATCCCA	4626
927	AACAUUUU G UACAAAAA	1604	TTTTGTGA GGCTAGCTACAACGA AATATGTT	4627
944	AUCAAAAU G UGUUUUAG	1605	CTAAACA GGCTAGCTACAACGA ATTTTGAT	4628
946	CAAAUGU G UUUUAGGA	1606	TCCTAAAA GGCTAGCTACAACGA ACATTTTG	4629
963	AACUUCU G UAAACAGG	1607	CCTGTTTA GGCTAGCTACAACGA AGGAAGTT	4630
991	GAAAGUAU G UCAACGAA	1608	TTGCTTGA GGCTAGCTACAACGA ATACTTTC	4631
1002	AACGAAUU G UGGGUCUU	1609	AAGACCCA GGCTAGCTACAACGA AATTCGTT	4632
1039	CACGCAU G UGGAUUAU	1610	AATATCCA GGCTAGCTACAACGA ATTGCGTG	4633
1137	AACAGUAU G UGAACCUU	1611	AAGGTTCA GGCTAGCTACAACGA ATACTGTT	4634
1184	UGCCAAGU G UUUGCUGA	1612	TCAGCAA GGCTAGCTACAACGA ACTTGGCA	4635
1251	GAACUUU G UGUCUCU	1613	AGGAGACA GGCTAGCTACAACGA AAAGGTTT	4636
1253	ACCUUUGU G UCUCUCU	1614	AGAGGAGA GGCTAGCTACAACGA ACAAGGT	4637
1294	AGCCGCUU G UUUUGCUC	1615	GAGCAAAA GGCTAGCTACAACGA AAGCGGCT	4638
1344	ACAAUUCU G UCGUGCUC	1616	GAGCACGA GGCTAGCTACAACGA AGAATTGT	4639
1390	GCUAGGCU G UGCUGCCA	1617	TGCGAGCA GGCTAGCTACAACGA AGCCTAGC	4640
1425	CGUCCUUU G UUUACGUC	1618	GACGTAA GGCTAGCTACAACGA AAAGGACG	4641
1508	CGCCUAU G UACCGACC	1619	GGTCGGTA GGCTAGCTACAACGA AATAGGCG	4642
1557	CCCCGUCU G UGCCUUCU	1620	AGAAGGCA GGCTAGCTACAACGA AGACGGGG	4643
1581	CGGACCGU G UGCACUUC	1621	GAAGTGCA GGCTAGCTACAACGA ACGGTCCG	4644
1684	UCAGCAAU G UCAACGAC	1622	GTCGTTGA GGCTAGCTACAACGA ATTGCTGA	4645
1719	CAAAGACU G UGUGUUUA	1623	TAAACACA GGCTAGCTACAACGA AGTCTTTG	4646
1721	AAGACUGU G UGUUUAU	1624	ATTAAACA GGCTAGCTACAACGA ACAGTCTT	4647
1723	GACUGUGU G UUUAAUGA	1625	TCATTAAA GGCTAGCTACAACGA ACACAGTC	4648
1772	AGGCUUUU G UACUAGGA	1626	TCCTAGTA GGCTAGCTACAACGA AAAGACCT	4649
1785	AGGAGGCU G UAGGCAUA	1627	TATGCCTA GGCTAGCTACAACGA AGCCTCCT	4650
1801	AAAUUGGU G UGUUCACC	1628	GGTGAACA GGCTAGCTACAACGA ACCAATTT	4651
1803	AUUGUGUGU G UUCACCAG	1629	CTGGTGAA GGCTAGCTACAACGA ACACCAAT	4652
1850	CAUCUCAU G UUCAUGUC	1630	GACATGAA GGCTAGCTACAACGA ATGAGATG	4653

Table 41

1856	AUGUUCAU G UCCUACUG	1631	CAGTAGGA GGCTAGCTACAACGA ATGAACAT	4654
1864	GUCCUACU G UUCAAGCC	1632	GGCTTGAA GGCTAGCTACAACGA AGTAGGAC	4655
1881	UCCAAGCU G UGCCUUGG	1633	CCAAGGCA GGCTAGCTACAACGA AGCTTGGA	4656
1939	GAGCUUCU G UGGAGUUA	1634	TAACTCCA GGCTAGCTACAACGA AGAAGCTC	4657
2013	UCUGCUCU G UAUCGGGG	1635	CCCCGATA GGCTAGCTACAACGA AGAGCAGA	4658
2045	GGAACAUU G UUCACCUC	1636	GAGGTGAA GGCTAGCTACAACGA AATGTTCC	4659
2082	GCUAUUCU G UGUUGGGG	1637	CCCCAACA GGCTAGCTACAACGA AGAATAGC	4660
2084	UAUUCUGU G UUGGGGUG	1638	CACCCCAA GGCTAGCTACAACGA ACAGAAATA	4661
2167	UCAGCUAU G UCAACGUU	1639	AACGTTGA GGCTAGCTACAACGA ATAGCTGA	4662
2205	CAACUAUU G UGGUUUCA	1640	TGAAACCA GGCTAGCTACAACGA AATAGTTG	4663
2222	CAUUUCCU G UCUUACUU	1641	AAGTAAGA GGCTAGCTACAACGA AGGAAATG	4664
2245	GAGAAACU G UUCUUGAA	1642	TTCAAGAA GGCTAGCTACAACGA AGTTTCTC	4665
2262	UAUUUGGU G UCUUUUGG	1643	CCAAAAGA GGCTAGCTACAACGA ACCAAATA	4666
2274	UUUGGAGU G UGGAUUCG	1644	CGAATCCA GGCTAGCTACAACGA ACTCCAAA	4667
2344	AAACUACU G UUGUUAGA	1645	TCTAACAA GGCTAGCTACAACGA AGTAGTTT	4668
2347	CUACUGUU G UUAGACGA	1646	TCGTCTAA GGCTAGCTACAACGA AACAGTAG	4669
2450	AUCUCAAU G UUAGUAUU	1647	AATACTAA GGCTAGCTACAACGA ATTGAGAT	4670
2573	AGGACAUU G UUGUAAGA	1648	TCTATCAA GGCTAGCTACAACGA AATGTCTT	4671
2583	UGAUAGAU G UAAGCAAU	1649	ATTGCTTA GGCTAGCTACAACGA ATCTATCA	4672
2594	AGCAAUUU G UGGGGCCC	1650	GGGCCCCA GGCTAGCTACAACGA AAATTGCT	4673
2663	AUCCCAAU G UUACUAAA	1651	TTTAGTAA GGCTAGCTACAACGA ATTGGGAT	4674
2717	CAGAGUAU G UAGUAAU	1652	ATTAATA GGCTAGCTACAACGA ATACTCTG	4675
2901	AUCUUUCU G UCCCCAAU	1653	ATTGGGGA GGCTAGCTACAACGA AGAAAGAT	4676
3071	GGGGGACU G UUGGGGUG	1654	CACCCCAA GGCTAGCTACAACGA AGTCCCCC	4677
3111	UCACAACU G UGCCAGCA	1655	TGCTGGCA GGCTAGCTACAACGA AGTTGTGA	4678
40	AUCCCAGA G UCAGGGCC	1656	GGCCCTGA GGCTAGCTACAACGA TCTGGGAT	4679
46	GAGUCAGG G CCCUGUAC	1657	GTACAGGG GGCTAGCTACAACGA CCTGACTC	4680
65	UCCUGCUG G UGGCUCCA	1658	TGGAGCCA GGCTAGCTACAACGA CAGCAGGA	4681
68	UGCUGGUG G CUCCAGUU	1659	AACTGGAG GGCTAGCTACAACGA CACCAGCA	4682
74	UGGCUCCA G UUCAGGAA	1660	TTCTTGAA GGCTAGCTACAACGA TGGAGCCA	4683
85	CAGGAACA G UGAGCCCU	1661	AGGGCTCA GGCTAGCTACAACGA TGTTCCTG	4684
89	AACAGUGA G CCCUGCUC	1662	GAGCAGGG GGCTAGCTACAACGA TCACTGTT	4685
120	GCCAUAUC G UCAAUCUU	1663	AAGATTGA GGCTAGCTACAACGA GATATGGC	4686
196	CCCUGCUC G UGUUACAG	1664	CTGTAAAC GGCTAGCTACAACGA GAGCAGGG	4687
205	UGUUACAG G CGGGGUUU	1665	AAACCCCG GGCTAGCTACAACGA CTGTAAAC	4688
210	CAGGCGGG G UUUUUCUU	1666	AAGAAAAA GGCTAGCTACAACGA CCCGCTG	4689
248	ACCACAGA G UCUAGACU	1667	AGTCTAGA GGCTAGCTACAACGA TCTGTGGT	4690
258	CUAGACUC G UGGUGGAC	1668	GTCCACCA GGCTAGCTACAACGA GAGTCTAG	4691
261	GACUCGUG G UGGACUUC	1669	GAAGTCCA GGCTAGCTACAACGA CACGAGTC	4692
295	GAACACCC G UGUGUCUU	1670	AAGACACA GGCTAGCTACAACGA GGGTGTTC	4693
305	GUGUCUUG G CCAAAAUU	1671	AATTTTGG GGCTAGCTACAACGA CAAGACAC	4694
318	AAUUCGCA G UCCCAAU	1672	ATTTGGGA GGCTAGCTACAACGA TGCGAATT	4695
332	AAUCUCCA G UCACUCAC	1673	GTGAGTGA GGCTAGCTACAACGA TGGAGATT	4696
368	UUGUCCUG G UUAUCGCU	1674	AGCGATAA GGCTAGCTACAACGA CAGGACAA	4697
390	UGUCGCG G CGUUUUAU	1675	ATAAAACG GGCTAGCTACAACGA CGCAGACA	4698
392	UCUGCGGC G UUUUAUCA	1676	TGATAAAA GGCTAGCTACAACGA GCCGCAGA	4699
442	UCUUGUUG G UUCUUCUG	1677	CAGAGAAA GGCTAGCTACAACGA CAACAAGA	4700
461	CUAUCUAG G UAUGUUGC	1678	GCAACATA GGCTAGCTACAACGA CTTGATAG	4701
472	UGUUGCCC G UUGUCCU	1679	AGGACAAA GGCTAGCTACAACGA GGGCAACA	4702
506	AACAACCA G CACCGGAC	1680	GTCCGCTG GGCTAGCTACAACGA TGGTTGTT	4703
625	CAUCUUGG G CUUUCGCA	1681	TGCGAAAG GGCTAGCTACAACGA CCAAGATG	4704

Table 41

648	CUAUGGGA G UGGGCCUC	1682	GAGGCCCA GGCTAGCTACAACGA TCCCATAG	4705
652	GGGAGUGG G CCUCAGUC	1683	GACTGAGG GGCTAGCTACAACGA CCACTCCC	4706
658	GGGCCUCA G UCCGUUUC	1684	GAAACGGA GGCTAGCTACAACGA TGAGGCCC	4707
662	CUCAGUCC G UUUCUCUU	1685	AAGAGAAA GGCTAGCTACAACGA GGACTGAG	4708
672	UUCUCUUG G CUCAGUUU	1686	AAACTGAG GGCTAGCTACAACGA CAAGAGAA	4709
677	UUGGCUCA G UUUACUAG	1687	CTAGTAAA GGCTAGCTACAACGA TGAGCCAA	4710
685	GUUUACUA G UGCCAUUU	1688	AAATGGCA GGCTAGCTACAACGA TAGTAAAC	4711
699	UUUGUUCA G UGGUUCGU	1689	ACGAACCA GGCTAGCTACAACGA TGAACAAA	4712
702	GUUCAGUG G UUCGUAGG	1690	CCTACGAA GGCTAGCTACAACGA CACTGAAC	4713
706	AGUGGUUC G UAGGGCUU	1691	AAGCCCTA GGCTAGCTACAACGA GAACCACT	4714
711	UUCGUAGG G CUUUCCCC	1692	GGGGAAAG GGCTAGCTACAACGA CCTACGAA	4715
729	ACUGUCUG G CUUUCAGU	1693	ACTGAAAG GGCTAGCTACAACGA CAGACAGT	4716
736	GGCUUUCA G UUAUAUGG	1694	CCATATAA GGCTAGCTACAACGA TGAAAGCC	4717
753	AUGAUGUG G UUUGGGG	1695	CCCCAAAA GGCTAGCTACAACGA CACATCAT	4718
762	UUUUGGGG G CCAAGUCU	1696	AGACTTGG GGCTAGCTACAACGA CCCCCAAA	4719
767	GGGGCCAA G UCUGUACA	1697	TGTACAGA GGCTAGCTACAACGA TTGGCCCC	4720
785	CAUCUUGA G UCCCUUUA	1698	TAAAGGGA GGCTAGCTACAACGA TCAAGATG	4721
826	GUCUUUGG G UAUACAUU	1699	AATGTATA GGCTAGCTACAACGA CCAAAGAC	4722
898	AAUUGGGA G UUGGGGCA	1700	TGCCCCAA GGCTAGCTACAACGA TCCCAATT	4723
904	GAGUUGGG G CACAUUGC	1701	GCAATGTG GGCTAGCTACAACGA CCAACTC	4724
971	GUAAACAG G CCUAUUGA	1702	TCAATAGG GGCTAGCTACAACGA CTGTTTAC	4725
987	AUUGGAAA G UAUGUCAA	1703	TTGACATA GGCTAGCTACAACGA TTTCCAAT	4726
1006	AAUUGUGG G UCUUUUGG	1704	CCAAAAGA GGCTAGCTACAACGA CCACAATT	4727
1016	CUUUUGGG G UUUGCCGC	1705	GCGGCAAA GGCTAGCTACAACGA CCAAAAG	4728
1080	GCAUACAA G CAAAACAG	1706	CTGTTTTG GGCTAGCTACAACGA TTGTATGC	4729
1089	CAAAACAG G CUUUUACU	1707	AGTAAAAG GGCTAGCTACAACGA CTGTTTTG	4730
1116	CUUACAAG G CCUUUCUA	1708	TAGAAAGG GGCTAGCTACAACGA CTGTAAAG	4731
1126	CUUUCUAA G UAAACAGU	1709	ACTGTTTA GGCTAGCTACAACGA TTAGAAAG	4732
1133	AGUAAACA G UAUGUGAA	1710	TTCACATA GGCTAGCTACAACGA TGTTTACT	4733
1152	UUUACCCC G UUGCUCGG	1711	CCGAGCAA GGCTAGCTACAACGA GGGGTAAA	4734
1160	GUUGCUCG G CAACGGCC	1712	GGCCGTTG GGCTAGCTACAACGA CGAGCAAC	4735
1166	CGGCAACG G CCUGGUCU	1713	AGACCAGG GGCTAGCTACAACGA CGTTGCCG	4736
1171	ACGGCCUG G UCUAUGCC	1714	GGCATAGA GGCTAGCTACAACGA CAGGCCGT	4737
1182	UAUGCCAA G UGUUUGCU	1715	AGCAAACA GGCTAGCTACAACGA TTGGCATA	4738
1207	CCCCACUG G UUGGGGCU	1716	AGCCCCAA GGCTAGCTACAACGA CAGTGGGG	4739
1213	UGGUUGGG G CUUGGCCA	1717	TGGCCAAG GGCTAGCTACAACGA CCAACCA	4740
1218	GGGGCUUG G CCAUAGGC	1718	GCCTATGG GGCTAGCTACAACGA CAAGCCCC	4741
1225	GGCAUAG G CCAUCAGC	1719	GCTGATGG GGCTAGCTACAACGA CTATGGCC	4742
1232	GGCAUCA G CGCAUGCG	1720	CGCATGCG GGCTAGCTACAACGA TGATGGCC	4743
1240	GCGCAUGC G UGGAACCU	1721	AGGTTCCA GGCTAGCTACAACGA GCATGCGC	4744
1287	AACUCCUA G CCGCUUGU	1722	ACAAGCGG GGCTAGCTACAACGA TAGGAGTT	4745
1306	UGCUCGCA G CAGGUCUG	1723	CAGACCTG GGCTAGCTACAACGA TCGAGCA	4746
1310	CGCAGCAG G UCUGGGGC	1724	GCCCCAGA GGCTAGCTACAACGA CTGCTGCG	4747
1317	GGUCUGGG G CAAAACUC	1725	GAGTTTTG GGCTAGCTACAACGA CCCAGACC	4748
1347	AUUCUGUC G UGCUCUCC	1726	GGAGAGCA GGCTAGCTACAACGA GACAGAAT	4749
1379	UUUCCAUG G CUGCUAGG	1727	CCTAGCAG GGCTAGCTACAACGA CATGGAAA	4750
1387	GCUGCUAG G CUGUGCUG	1728	CAGCACAG GGCTAGCTACAACGA CTAGCAGC	4751
1418	CGCGGGAC G UCCUUGU	1729	ACAAAGGA GGCTAGCTACAACGA GTCCCGCG	4752
1431	UUGUUUAC G UCCGUCG	1730	CGACGGGA GGCTAGCTACAACGA GTAAACAA	4753
1436	UACGUCCC G UCGGCGCU	1731	AGCGCGGA GGCTAGCTACAACGA GGGACGTA	4754
1440	UCCCGUCG G CGCUGAAU	1732	ATTCAGCG GGCTAGCTACAACGA CGACGGGA	4755

Table 41

1471	CUCCCCGG G CCGCUUGG	1733	CCAAGCGG GGCTAGCTACAACGA CCCGGGAG	4756
1481	CGCUUGGG G CUCUACCG	1734	CGGTAGAG GGCTAGCTACAACGA CCAAGCG	4757
1517	UACCGACC G UCCACGGG	1735	CCCGTGGA GGCTAGCTACAACGA GGTCGGTA	4758
1526	UCCACGGG G CGCACCUC	1736	GAGGTGCG GGCTAGCTACAACGA CCGTGGA	4759
1553	GACUCCCC G UCUGUGCC	1737	GGCACAGA GGCTAGCTACAACGA GGGGAGTC	4760
1579	GCCGGACC G UGUGCACU	1738	AGTGCACA GGCTAGCTACAACGA GTCCGGC	4761
1605	CUCUGCAC G UCGCAUGG	1739	CCATGCGA GGCTAGCTACAACGA GTGCAGAG	4762
1622	AGACCACC G UGAACGCC	1740	GGCGTTCA GGCTAGCTACAACGA GGTGGTCT	4763
1649	UGCCCAAG G UCUUGCAU	1741	ATGCAAGA GGCTAGCTACAACGA CTTGGGCA	4764
1679	GACUUUCA G CAAUGUCA	1742	TGACATTG GGCTAGCTACAACGA TGAAGTC	4765
1703	ACCUGAG G CAUACUUC	1743	GAAGTATG GGCTAGCTACAACGA CTAAGGT	4766
1732	UUUAAUGA G UGGGAGGA	1744	TCCTCCCA GGCTAGCTACAACGA TCATTAA	4767
1741	UGGGAGGA G UUGGGGA	1745	TCCCCCAA GGCTAGCTACAACGA TCCTCCCA	4768
1754	GGGAGGAG G UUAGGUUA	1746	TAACCTAA GGCTAGCTACAACGA CTCCTCCC	4769
1759	GAGGUUAG G UUAAGGU	1747	ACCTTTAA GGCTAGCTACAACGA CTAACCTC	4770
1766	GGUUAAG G UCUUUGUA	1748	TACAAAGA GGCTAGCTACAACGA CTTTAACC	4771
1782	ACUAGGAG G CUGUAGGC	1749	GCCTACAG GGCTAGCTACAACGA CTCCTAGT	4772
1789	GGCUGUAG G CAUAAAUU	1750	AATTTATG GGCTAGCTACAACGA CTACAGCC	4773
1799	AUAAAUUG G UGUGUUA	1751	TGAACACA GGCTAGCTACAACGA CAATTTAT	4774
1811	GUUCACCA G CACCAUGC	1752	GCATGGTG GGCTAGCTACAACGA TGGTGAAC	4775
1870	CUGUUCAA G CCUCCAAG	1753	CTTGAGG GGCTAGCTACAACGA TTGAACAG	4776
1878	GCCUCCAA G CUGUGCCU	1754	AGGCACAG GGCTAGCTACAACGA TTGGAGGC	4777
1890	UGCCUUGG G UGGCUUUG	1755	CAAAGCCA GGCTAGCTACAACGA CCAAGGCA	4778
1893	CUUGGGUG G CUUUGGGG	1756	CCCCAAG GGCTAGCTACAACGA CACCCAAG	4779
1901	GCUUUGGG G CAUGGACA	1757	TGTCCATG GGCTAGCTACAACGA CCCAAGC	4780
1917	AUUGACCC G UAUAAAGA	1758	TCTTTATA GGCTAGCTACAACGA GGGTCAAT	4781
1933	AAUUUGGA G CUUCUGUG	1759	CACAGAAG GGCTAGCTACAACGA TCCAAATT	4782
1944	UCUGUGGA G UUACUCUC	1760	GAGAGTAA GGCTAGCTACAACGA TCCACAGA	4783
2023	AUCGGGGG G CCUAGAG	1761	CTCTAAGG GGCTAGCTACAACGA CCCCCGAT	4784
2031	GCCUUAGA G UCUCGGA	1762	TCCGGAGA GGCTAGCTACAACGA TCTAAGGC	4785
2062	ACCAUACG G CACUCAGG	1763	CCTGAGTG GGCTAGCTACAACGA CGTATGGT	4786
2070	GCACUCAG G CAAGCUAU	1764	ATAGCTTG GGCTAGCTACAACGA CTGAGTGC	4787
2074	UCAGGCAA G CUAUUCUG	1765	CAGAATAG GGCTAGCTACAACGA TTGCCTGA	4788
2090	GUGUUGGG G UGAGUUGA	1766	TCAACTCA GGCTAGCTACAACGA CCAACAC	4789
2094	UGGGUGA G UUGAUGAA	1767	TTCATCAA GGCTAGCTACAACGA TCACCCA	4790
2107	UGAAUCUA G CCACCGG	1768	CCAGGTGG GGCTAGCTACAACGA TAGATTCA	4791
2116	CCACCGG G UGGGAAGU	1769	ACTTCCCA GGCTAGCTACAACGA CCAGGTGG	4792
2123	GGUGGGAA G UAAUUUGG	1770	CCAAATTA GGCTAGCTACAACGA TTCCACC	4793
2140	AAGAUCCA G CAUCCAGG	1771	CCTGGATG GGCTAGCTACAACGA TGGATCTT	4794
2155	GGGAUUUA G UAGUCAGC	1772	GCTGACTA GGCTAGCTACAACGA TAATTCCC	4795
2158	AAUUGUA G UCAGCUAU	1773	ATAGCTGA GGCTAGCTACAACGA TACTAATT	4796
2162	AGUAGUCA G CUAUGUCA	1774	TGACATAG GGCTAGCTACAACGA TGACTACT	4797
2173	AUGUCAAC G UUAUAUG	1775	CATATTAA GGCTAGCTACAACGA GTTGACAT	4798
2183	UAAUUGG G CCUAAAA	1776	TTTTTAGG GGCTAGCTACAACGA CCATATTA	4799
2208	CUAUUGUG G UUCACAU	1777	ATGTGAAA GGCTAGCTACAACGA CACAATAG	4800
2235	ACUUUUGG G CGAGAAAC	1778	GTTTCTCG GGCTAGCTACAACGA CCAAAAGT	4801
2260	AAUAUUUG G UGUUUUU	1779	AAAAGACA GGCTAGCTACAACGA CAAATATT	4802
2272	CUUUUGGA G UGUGGAUU	1780	AATCCACA GGCTAGCTACAACGA TCCAAAAG	4803
2360	ACGAAGAG G CAGGUCCC	1781	GGGACCTG GGCTAGCTACAACGA CTCTTCGT	4804
2364	AGAGGCAG G UCCCCUAG	1782	CTAGGGGA GGCTAGCTACAACGA CTGCCTCT	4805
2403	AGACGAAG G UCUCAAUC	1783	GATTGAGA GGCTAGCTACAACGA CTTGCTCT	4806

Table 41

2417	AUCGCCG G UCGCAGAA	1784	TTCTGCGA GGCTAGCTACAACGA GCGGCGAT	4807
2454	CAAUGUUA G UAUUCCUU	1785	AAGGAATA GGCTAGCTACAACGA TAACATTG	4808
2474	CACAUAA G UGGGAAAC	1786	GTTTCCCA GGCTAGCTACAACGA CTTATGTG	4809
2491	UUUACGGG G CUUUUUAUC	1787	GAATAAAG GGCTAGCTACAACGA CCCGTAAA	4810
2507	CUUCUACG G UACCUUGC	1788	GCAAGGTA GGCTAGCTACAACGA CGTAGAAG	4811
2530	CCUAAAUG G CAAACUCC	1789	GGAGTTTG GGCTAGCTACAACGA CATTTAGG	4812
2587	AGAUGUAA G CAAUUGU	1790	ACAAATTG GGCTAGCTACAACGA TTACATCT	4813
2599	UUUGUGGG G CCCCUUAC	1791	GTAAGGGG GGCTAGCTACAACGA CCCACAAA	4814
2609	CCCUUACA G UAAAUGAA	1792	TTCATTTA GGCTAGCTACAACGA TGTAAGGG	4815
2650	CCUGCUAG G UUUUAUCC	1793	GGATAAAA GGCTAGCTACAACGA CTAGCAGG	4816
2701	AUCAAAAC G UAUUAUCC	1794	GGATAATA GGCTAGCTACAACGA GGTTTGAT	4817
2713	UAUCCAGA G UAUGUAGU	1795	ACTACATA GGCTAGCTACAACGA TCTGGATA	4818
2720	AGUAUGUA G UUAUUAU	1796	ATGATTAA GGCTAGCTACAACGA TACATACT	4819
2768	UUUGGAAG G CGGGGAUC	1797	GATCCCCG GGCTAGCTACAACGA CTTCCAAA	4820
2791	AAAAGAGA G UCCACACG	1798	CGTGTGGA GGCTAGCTACAACGA TCTCTTTT	4821
2799	GUCCACAC G UAGCGCU	1799	AGGCGCTA GGCTAGCTACAACGA GTGTGGAC	4822
2802	CACACGUA G CGCCUCAU	1800	ATGAGGCG GGCTAGCTACAACGA TACGTGTG	4823
2818	UUUUGCGG G UCACCAUA	1801	TATGGTGA GGCTAGCTACAACGA CCGCAAAA	4824
2848	GAUCUACA G CAUGGGAG	1802	CTCCCATG GGCTAGCTACAACGA TGTAGATC	4825
2857	CAUGGGAG G UUGGUCUU	1803	AAGACCAA GGCTAGCTACAACGA CTCCCATG	4826
2861	GGAGGUUG G UCUUCCAA	1804	TTGGAAGA GGCTAGCTACAACGA CAACCTCC	4827
2881	UCGAAAAG G CAUGGGGA	1805	TCCCCATG GGCTAGCTACAACGA CTTTTCGA	4828
2936	GAUCAUCA G UUGGACCC	1806	GGGTCCAA GGCTAGCTACAACGA TGATGATC	4829
2955	CAUUCAAA G CCAACUCA	1807	TGAGTTGG GGCTAGCTACAACGA TTTGAATG	4830
2964	CCAACUCA G UAAAUCCA	1808	TGGATTTA GGCTAGCTACAACGA TGAGTTGG	4831
3005	GACAACUG G CCGGACGC	1809	GCGTCCGG GGCTAGCTACAACGA CAGTTGTC	4832
3021	CCAACAAG G UGGGAGUG	1810	CACTCCCA GGCTAGCTACAACGA CTTGTTGG	4833
3027	AGGUGGGA G UGGGAGCA	1811	TGCTCCCA GGCTAGCTACAACGA TCCCACCT	4834
3033	GAGUGGGA G CAUUCGGG	1812	CCCGAATG GGCTAGCTACAACGA TCCCCTC	4835
3041	GCAUUCGG G CCAGGGUU	1813	AACCCTGG GGCTAGCTACAACGA CCGAATGC	4836
3047	GGGCCAGG G UUCACCCC	1814	GGGGTGAA GGCTAGCTACAACGA CCTGGCCC	4837
3077	CUGUUGGG G UGGAGCCC	1815	GGGCTCCA GGCTAGCTACAACGA CCAACAG	4838
3082	GGGUGGGA G CCCUCACG	1816	CGTGAGGG GGCTAGCTACAACGA TCCACCCC	4839
3097	CGCUCAGG G CCUACUCA	1817	TGAGTAGG GGCTAGCTACAACGA CCTGAGCG	4840
3117	CUGUGCCA G CAGUCCU	1818	AGGAGCTG GGCTAGCTACAACGA TGGCACAG	4841
3120	UGCCAGCA G CUCCUCCU	1819	AGGAGGAG GGCTAGCTACAACGA TGCTGGCA	4842
3146	ACCAAUCG G CAGUCAGG	1820	CCTGACTG GGCTAGCTACAACGA CGATTGGT	4843
3149	AAUCGGCA G UCAGGAAG	1821	CTTCCTGA GGCTAGCTACAACGA TGCCGATT	4844
3158	UCAGGAAG G CAGCCUAC	1822	GTAGGCTG GGCTAGCTACAACGA CTTCCTGA	4845
3161	GGAAGGCA G CCUACUCC	1823	GGAGTAGG GGCTAGCTACAACGA TGCCCTCC	4846
3204	AUCCUCAG G CCAUGCAG	1824	CTGCATGG GGCTAGCTACAACGA CTGAGGAT	4847
10	ACUCCACC A CUUCCAC	1825	GTGGAAAG GGCTAGCTACAACGA GGTGGAGT	4848
17	CACUUUCC A CCAAACUC	1826	GAGTTTGG GGCTAGCTACAACGA GGAAAGTG	4849
22	UCCACCAA A CUCUCAA	1827	TTGAAGAG GGCTAGCTACAACGA TTGGTGGA	4850
32	UCUUAAG A UCCAGAG	1828	CTCTGGGA GGCTAGCTACAACGA CTTGAAGA	4851
53	GGCCUGU A CUUUCUG	1829	CAGGAAAG GGCTAGCTACAACGA ACAGGGCC	4852
82	GUUCAGGA A CAGUGAGC	1830	GCTCACTG GGCTAGCTACAACGA TCCTGAAC	4853
101	UGCUCAGA A UACUGUCU	1831	AGACAGTA GGCTAGCTACAACGA TCTGAGCA	4854
103	CUCAGAAU A CUGUCUCU	1832	AGAGACAG GGCTAGCTACAACGA ATTCTGAG	4855
115	UCUCUGCC A UAUCGUCA	1833	TGACGATA GGCTAGCTACAACGA GGCAGAGA	4856
117	UCUGCCAU A UCGUCAU	1834	ATTGACGA GGCTAGCTACAACGA ATGGCAGA	4857

Table 4 I

124	UAUCGUCA A UCUAUCG	1835	CGATAAGA GGCTAGCTACAACGA TGACGATA	4858
129	UCAAUCUU A UCGAAGAC	1836	GTCTTCGA GGCTAGCTACAACGA AAGATTGA	4859
136	UAUCGAAG A CUGGGGAC	1837	GTCCCCAG GGCTAGCTACAACGA CTTCGATA	4860
143	GACUGGGG A CCCUGUAC	1838	GTACAGGG GGCTAGCTACAACGA CCCAGTC	4861
150	GACCCUGU A CCGAACAU	1839	ATGTTCGG GGCTAGCTACAACGA ACAGGGTC	4862
155	UGUACCGA A CAUGGAGA	1840	TCTCCATG GGCTAGCTACAACGA TCGGTACA	4863
157	UACCGAAC A UGGAGAAC	1841	GTTCTCCA GGCTAGCTACAACGA GTTCGGTA	4864
164	CAUGGAGA A CAUCGCAU	1842	ATGCGATG GGCTAGCTACAACGA TCTCCATG	4865
166	UGGAGAAC A UCGCAUCA	1843	TGATGCGA GGCTAGCTACAACGA GTTCTCCA	4866
171	AACAUCGC A UCAGGACU	1844	AGTCTCTG GGCTAGCTACAACGA GCGATGTT	4867
177	GCAUCAGG A CUCCUAGG	1845	CCTAGGAG GGCTAGCTACAACGA CCTGATGC	4868
186	CUCCUAGG A CCCUGGCU	1846	AGCAGGGG GGCTAGCTACAACGA CCTAGGAG	4869
201	CUCGUGUU A CAGGCGGG	1847	CCCGCCTG GGCTAGCTACAACGA AACACGAG	4870
223	UCUUGUUG A CAAAAAUC	1848	GATTTTGT GGCTAGCTACAACGA CAACAAGA	4871
229	UGACAAA A UCCUCACA	1849	TGTGAGGA GGCTAGCTACAACGA TTTGTCTA	4872
235	AAAUCCUC A CAUACCA	1850	TGGTATTG GGCTAGCTACAACGA GAGGATTT	4873
238	UCCUCACA A UACCACAG	1851	CTGTGGTA GGCTAGCTACAACGA TGTGAGGA	4874
240	CUCACAAU A CCACAGAG	1852	CTCTGTGG GGCTAGCTACAACGA ATTGTGAG	4875
243	ACAAUACC A CAGAGUCU	1853	AGACTCTG GGCTAGCTACAACGA GGTATTGT	4876
254	GAGUCUAG A CUCGUGGU	1854	ACCACGAG GGCTAGCTACAACGA CTAGACTC	4877
265	CGUGGUGG A CUUCUCUC	1855	GAGAGAAG GGCTAGCTACAACGA CCACCACG	4878
275	UUCUCUCA A UUUUCUAG	1856	CTAGAAAA GGCTAGCTACAACGA TGAGAGAA	4879
289	UAGGGGGA A CACCCGUG	1857	CACGGGTG GGCTAGCTACAACGA TCCCCCTA	4880
291	GGGGGAAC A CCCGUGUG	1858	CACACGGG GGCTAGCTACAACGA GTTCCCCC	4881
311	UGGCCAAA A UUCGCAGU	1859	ACTGCGAA GGCTAGCTACAACGA TTTGGCCA	4882
325	AGUCCCAA A UCUCAGU	1860	ACTGGAGA GGCTAGCTACAACGA TTGGGACT	4883
335	CUCCAGUC A CUCACCAA	1861	TTGGTGAG GGCTAGCTACAACGA GACTGGAG	4884
339	AGUCACUC A CCAACCUG	1862	CAGGTTGG GGCTAGCTACAACGA GAGTGAAT	4885
343	ACUCACCA A CCUGUUGU	1863	ACAACAGG GGCTAGCTACAACGA TGGTGAGT	4886
358	GUCCUCCA A UUUGUCCU	1864	AGGACAAA GGCTAGCTACAACGA TGGAGGAC	4887
371	UCUGGUU A UCGCUGGA	1865	TCCAGCGA GGCTAGCTACAACGA AACCAGGA	4888
379	AUCGUGG A UGUGUCUG	1866	CAGACACA GGCTAGCTACAACGA CCAGCGAT	4889
397	GGCGUUU A UCAUCUUC	1867	GAAGATGA GGCTAGCTACAACGA AAAACGCC	4890
400	GUUUUAUC A UCUCUCC	1868	GAGGAAGA GGCTAGCTACAACGA GATAAAAC	4891
412	UCCUCUGC A UCCUGCUG	1869	CAGCAGGA GGCTAGCTACAACGA GCAGAGGA	4892
423	CUGCUGCU A UGCCUCAU	1870	ATGAGGCA GGCTAGCTACAACGA AGCAGCAG	4893
430	UAUGCCUC A UCUCUUG	1871	CAAGAAGA GGCTAGCTACAACGA GAGGCATA	4894
452	UCUCUGG A CUAUCAAG	1872	CTTGATAG GGCTAGCTACAACGA CCAGAAGA	4895
455	UCUGGACU A UCAAGGUA	1873	TACCTTGA GGCTAGCTACAACGA AGTCCAGA	4896
463	AUCAAGGU A UGUUGCCC	1874	GGGCAACA GGCTAGCTACAACGA ACCTTGAT	4897
484	GUCCUCUA A UUCAGGA	1875	TCCTGGAA GGCTAGCTACAACGA TAGAGGAC	4898
492	AUUCAGG A UCAUCAAC	1876	GTTGATGA GGCTAGCTACAACGA CCTGGAAT	4899
495	CCAGGAUC A UCAACAAC	1877	GTTGTTGA GGCTAGCTACAACGA GATCCTGG	4900
499	GAUCAUCA A CAACGAGC	1878	GCTGGTTG GGCTAGCTACAACGA TGATGATC	4901
502	CAUCAACA A CCAGCACC	1879	GGTGCTGG GGCTAGCTACAACGA TGTGATG	4902
513	AGCACCGG A CCAUGCAA	1880	TTGCAATG GGCTAGCTACAACGA CCGGTGCT	4903
516	ACCGGACC A UGCAAAAC	1881	GTTTTCGA GGCTAGCTACAACGA GGTCCGGT	4904
523	CAUGCAAA A CCUGCACA	1882	TGTGCAAG GGCTAGCTACAACGA TTTGCATG	4905
529	AAACCUGC A CAACUCCU	1883	AGGAGTTG GGCTAGCTACAACGA GCAGGTTT	4906
532	CCUGCACA A CUCCUGCU	1884	AGCAGGAG GGCTAGCTACAACGA TGTGCAGG	4907
547	CUCAAGGA A CCUCUAUG	1885	CATAGAGG GGCTAGCTACAACGA TCCTTGAG	4908

Table 41

553	GAACCUCU A UGUUUCCC	1886	GGGAAACA GGCTAGCTACAACGA AGAGGTTC	4909
564	UUUCCUC A UGUUGCUG	1887	CAGCAACA GGCTAGCTACAACGA GAGGGAAA	4910
574	GUUGCUGU A CAAAACCU	1888	AGGTTTTG GGCTAGCTACAACGA ACAGCAAC	4911
579	UGUACAAA A CCUACGGA	1889	TCCGTAGG GGCTAGCTACAACGA TTTGTACA	4912
583	CAAAACCU A CGGACGGA	1890	TCCGTCCG GGCTAGCTACAACGA AGGTTTTG	4913
587	ACCUCAGG A CGGAAACU	1891	AGTTTCCG GGCTAGCTACAACGA CCGTAGGT	4914
593	GGACGGAA A CUGCACCU	1892	AGGTGCAG GGCTAGCTACAACGA TTCGTCC	4915
598	GAAACUGC A CCUGUAU	1893	AATACAGG GGCTAGCTACAACGA GCAGTTTC	4916
604	GCACUGU A UUCCCAUC	1894	GATGGGAA GGCTAGCTACAACGA ACAGGTGC	4917
610	GUUUUCCC A UCCCAUCA	1895	TGATGGGA GGCTAGCTACAACGA GGAATAC	4918
615	CCCAUCCC A UCAUCUUG	1896	CAAGATGA GGCTAGCTACAACGA GGGATGGG	4919
618	AUCCCAUC A UCUUGGGC	1897	GCCCAAGA GGCTAGCTACAACGA GATGGGAT	4920
636	UUCGCAAA A UACCUAUG	1898	CATAGGTA GGCTAGCTACAACGA TTTGCGAA	4921
638	CGCAAAAU A CCUAUGGG	1899	CCCATAGG GGCTAGCTACAACGA ATTTGCG	4922
642	AAAUACCU A UGGGAGUG	1900	CACTCCCA GGCTAGCTACAACGA AGGTATTT	4923
681	CUCAGUUU A CUAGUGCC	1901	GGCACTAG GGCTAGCTACAACGA AACTGAG	4924
690	CUAGUGCC A UUUGUUA	1902	TGAACAAA GGCTAGCTACAACGA GGCACTAG	4925
721	UUUCCCCC A CUGUCUGG	1903	CCAGACAG GGCTAGCTACAACGA GGGGAAA	4926
739	UUUCAGUU A UAUGGAUG	1904	CATCCATA GGCTAGCTACAACGA AACTGAAA	4927
741	UCAGUUUA A UGGAUGAU	1905	ATCATCCA GGCTAGCTACAACGA ATAAGTGA	4928
745	UUUAUUGG A UGAUGUGG	1906	CCACATCA GGCTAGCTACAACGA CCATATA	4929
748	UAUGGAUG A UGUGGUUU	1907	AAACCACA GGCTAGCTACAACGA CATCCATA	4930
773	AAGUCUGU A CAACAUCU	1908	AGATGTTG GGCTAGCTACAACGA ACAGACTT	4931
776	UCUGUACA A CAUCUUGA	1909	TCAAGATG GGCTAGCTACAACGA TGTACAGA	4932
778	UGUACAAC A UCUUGAGU	1910	ACTCAAGA GGCTAGCTACAACGA GTTGTACA	4933
793	GUCCCUUU A UGCCGUG	1911	CAGCGGCA GGCTAGCTACAACGA AAAGGGAC	4934
804	CCGUGUUU A CCAUUUUU	1912	AAAATTGG GGCTAGCTACAACGA AACAGCGG	4935
808	UGUUACCA A UUUUCUUU	1913	AAAGAAAA GGCTAGCTACAACGA TGTAACA	4936
828	CUUUGGGU A UACAUUUA	1914	TAAATGTA GGCTAGCTACAACGA ACCCAAAG	4937
830	UUGGGUUA A CAUUUAAA	1915	TTTAAATG GGCTAGCTACAACGA ATACCCAA	4938
832	GGGUUAUC A UUUAAACC	1916	GGTTTAAA GGCTAGCTACAACGA GTATACCC	4939
838	ACAUUUAA A CCCUCACA	1917	TGTGAGGG GGCTAGCTACAACGA TTAATGT	4940
844	AAACCCUC A CAAAACAA	1918	TTGTTTTG GGCTAGCTACAACGA GAGGTTT	4941
849	CUCACAAA A CAAAAAGA	1919	TCITTTTG GGCTAGCTACAACGA TTTGTGAG	4942
857	ACAAAAAG A UGGGGAUA	1920	TATCCCCA GGCTAGCTACAACGA CTTTTGT	4943
863	AGAUGGGG A UAUUCCCU	1921	AGGGAATA GGCTAGCTACAACGA CCCCATCT	4944
865	AUGGGGAU A UUCCCUUA	1922	TAAGGGAA GGCTAGCTACAACGA ATCCCAT	4945
874	UUCCCUUA A CUUCAUGG	1923	CCATGAAG GGCTAGCTACAACGA TAAGGGAA	4946
879	UUAACUUC A UGGGAUUA	1924	ATATCCCA GGCTAGCTACAACGA GAAGTTAA	4947
884	UUCAUGGG A UAUGUAAU	1925	ATTACATA GGCTAGCTACAACGA CCCATGAA	4948
886	CAUGGGAU A UGUAAUUG	1926	CAATTACA GGCTAGCTACAACGA ATCCCATG	4949
891	GAUAUGUA A UUGGGAGU	1927	ACTCCCAA GGCTAGCTACAACGA TACATATC	4950
906	GUUGGGGC A CAUUGCCA	1928	TGGCAATG GGCTAGCTACAACGA GCCCAAC	4951
908	UGGGGCAC A UUGCCACA	1929	TGTGGCAA GGCTAGCTACAACGA GTGCCCA	4952
914	ACAUUGCC A CAGGAACA	1930	TGTTCCCTG GGCTAGCTACAACGA GGCAATGT	4953
920	CCACAGGA A CAUAUUGU	1931	ACAATATG GGCTAGCTACAACGA TCCTGTGG	4954
922	ACAGGAAC A UAUGUAC	1932	GTACAATA GGCTAGCTACAACGA GTTCTGT	4955
924	AGGAACAU A UUGUACAA	1933	TTGTACAA GGCTAGCTACAACGA ATGTTCT	4956
929	CAUAUUGU A CAAAAAU	1934	ATTTTTTG GGCTAGCTACAACGA ACAATATG	4957
936	UACAAAAA A UCAAAUUG	1935	CATTTTGA GGCTAGCTACAACGA TTTTGTGA	4958
942	AAAUCAAA A UGUGUUUU	1936	AAAACACA GGCTAGCTACAACGA TTTGATTT	4959

Table 41

956	UUUAGGAA A CUUCCUGU	1937	ACAGGAAG GGCTAGCTACAACGA TTCCTAAA	4960
967	UCCUGUAA A CAGGCCUA	1938	TAGGCCTG GGCTAGCTACAACGA TTACAGGA	4961
975	ACAGGCCU A UUGAUUGG	1939	CCAATCAA GGCTAGCTACAACGA AGGCCTGT	4962
979	GCCUAUUG A UUGGAAAG	1940	CTTTCCAA GGCTAGCTACAACGA CAATAGGC	4963
989	UGGAAAGU A UGUCAACG	1941	CGTTGACA GGCTAGCTACAACGA ACTTTCCA	4964
995	GU AUGUCA A CGAAUUGU	1942	ACAATTCTG GGCTAGCTACAACGA TGACATAC	4965
999	GUCAACGA A UUGUGGGU	1943	ACCCACAA GGCTAGCTACAACGA TCGTTGAC	4966
1032	CCCCUUUC A CGCAAUGU	1944	ACATTGCG GGCTAGCTACAACGA GAAAGGGG	4967
1037	UUCACGCA A UGUGGAUA	1945	TATCCACA GGCTAGCTACAACGA TCGGTGAA	4968
1043	CAAUGUGG A UAUUCUGC	1946	GCAGAATA GGCTAGCTACAACGA CCACATTG	4969
1045	AUGUGGAU A UUCUGCUU	1947	AAGCAGAA GGCTAGCTACAACGA ATCCACAT	4970
1056	CUGCUUUA A UGCCUUUA	1948	TAAAGGCA GGCTAGCTACAACGA TAAAGCAG	4971
1064	AUGCCUUU A UAUGCAUG	1949	CATGCATA GGCTAGCTACAACGA AAAGGCAT	4972
1066	GCCUUUAU A UGCAUGCA	1950	TGCATGCA GGCTAGCTACAACGA ATAAAGGC	4973
1070	UUUAUUGC A UGCAUACA	1951	TGTATGCA GGCTAGCTACAACGA GCATATAA	4974
1074	AUGCAUGC A UACAAGCA	1952	TGCTTGTA GGCTAGCTACAACGA GCATGCAT	4975
1076	GCAUGCAU A CAAGCAAA	1953	TTTGCTTG GGCTAGCTACAACGA ATGCATGC	4976
1085	CAAGCAAA A CAGGCUUU	1954	AAAGCCTG GGCTAGCTACAACGA TTTGCTTG	4977
1095	AGGCUUUU A CUUUCUCG	1955	CGAGAAAG GGCTAGCTACAACGA AAAAGCCT	4978
1107	UCUCGCCA A CUUACAAG	1956	CTTGTAAG GGCTAGCTACAACGA TGGCGAGA	4979
1111	GCCAACUU A CAAGCCU	1957	AGGCCTTG GGCTAGCTACAACGA AAGTTGGC	4980
1130	CUAAGUAA A CAGUAUGU	1958	ACATACTG GGCTAGCTACAACGA TTACTTAG	4981
1135	UAAACAGU A UGUGAACC	1959	GGTTCACA GGCTAGCTACAACGA ACTGTTTA	4982
1141	GU AUGUGA A CCUUUACC	1960	GGTAAAGG GGCTAGCTACAACGA TCACATAC	4983
1147	GAACCUUU A CCCCUGUG	1961	CAACGGGG GGCTAGCTACAACGA AAAGGTTT	4984
1163	GCUCGGCA A CGGCCUGG	1962	CCAGGCCG GGCTAGCTACAACGA TGCCGAGC	4985
1175	CCUGGUCU A UGCCAAGU	1963	ACTTGCCA GGCTAGCTACAACGA AGACCAGG	4986
1192	GUUUGCUG A CGCAACCC	1964	GGGTTGCG GGCTAGCTACAACGA CAGCAAAC	4987
1197	CUGACGCA A CCCCCACU	1965	AGTGGGGG GGCTAGCTACAACGA TCGGTCAG	4988
1203	CAACCCCC A CUGGUUGG	1966	CCAACCAG GGCTAGCTACAACGA GGGGGTTG	4989
1221	GCUUGGCC A UAGGCCAU	1967	ATGGCCTA GGCTAGCTACAACGA GGCCAAGC	4990
1228	CAUAGGCC A UCAGCGCA	1968	TGCGCTGA GGCTAGCTACAACGA GGCCTATG	4991
1236	AUCAGCGC A UGCGUGGA	1969	TCCACGCA GGCTAGCTACAACGA GCGCTGAT	4992
1245	UGCGUGGA A CCUUUGUG	1970	CACAAAGG GGCTAGCTACAACGA TCCACGCA	4993
1266	CUCUGCCG A UCCAUACC	1971	GGTATGGA GGCTAGCTACAACGA CGGCAGAG	4994
1270	GCCGAUCC A UACCGCGG	1972	CCGCGGTA GGCTAGCTACAACGA GGATCGGC	4995
1272	CGAUCCAU A CCGCGGAA	1973	TTCCGCGG GGCTAGCTACAACGA ATGGATCG	4996
1280	ACCGCGGA A CUCCUAGC	1974	GCTAGGAG GGCTAGCTACAACGA TCCGCGGT	4997
1322	GGGGCAAA A CUCAUCGG	1975	CCGATGAG GGCTAGCTACAACGA TTTGCCCC	4998
1326	CAAAACUC A UCGGGACU	1976	AGTCCCGA GGCTAGCTACAACGA GAGTTTTG	4999
1332	UCAUCGGG A CUGACAAU	1977	ATTGTCAG GGCTAGCTACAACGA CCCGATGA	5000
1336	CGGGACUG A CAAUUCUG	1978	CAGAATTG GGCTAGCTACAACGA CAGTCCCG	5001
1339	GACUGACA A UUCUGUCG	1979	CGACAGAA GGCTAGCTACAACGA TGTCAATC	5002
1361	UCCCGCAA A UAUACAUC	1980	GATGTATA GGCTAGCTACAACGA TTGCGGGA	5003
1363	CCGCAAAU A UACAUCAU	1981	ATGATGTA GGCTAGCTACAACGA ATTTGCGG	5004
1365	GCAAAUUA A CAUCAUUU	1982	AAATGATG GGCTAGCTACAACGA ATATTTGC	5005
1367	AAAUUAUC A UCAUUUCC	1983	GGAAATGA GGCTAGCTACAACGA GTATATTT	5006
1370	UAUACAUC A UUUCCAUG	1984	CATGGAAG GGCTAGCTACAACGA GATGTATA	5007
1376	UCAUUUCC A UGGCUGCU	1985	AGCAGCCA GGCTAGCTACAACGA GGAAATGA	5008
1399	UGCUGCCA A CUGGAUCC	1986	GGATCCAG GGCTAGCTACAACGA TGGCAGCA	5009
1404	CCAACUGG A UCCUACGC	1987	GCGTAGGA GGCTAGCTACAACGA CCAGTTGG	5010

Table 41

1409	UGGAUCCU A CGCGGGAC	1988	GTCCCCGG GGCTAGCTACAACGA AGGATCCA	5011
1416	UACGCGGG A CGUCCUUU	1989	AAAGGACG GGCTAGCTACAACGA CCCGCGTA	5012
1429	CUUUGUUU A CGUCCCGU	1990	ACGGGACG GGCTAGCTACAACGA AAACAAAG	5013
1447	GGGCUGA A UCCCGCGG	1991	CCGCGGGA GGCTAGCTACAACGA TCAGCGCC	5014
1456	UCCGCGG A CGACCCU	1992	AGGGGTG GGCTAGCTACAACGA CCGCGGGA	5015
1459	CGCGGACG A CCCUCCC	1993	GGGAGGGG GGCTAGCTACAACGA CGTCCGCG	5016
1486	GGGGCUCU A CCGCCCGC	1994	GCGGGCGG GGCTAGCTACAACGA AGAGCCCC	5017
1505	CUCCGCCU A UUGUACCG	1995	CGGTACAA GGCTAGCTACAACGA AGGCGGAG	5018
1510	CCUAUUGU A CCGACCGU	1996	ACGGTCGG GGCTAGCTACAACGA ACAATAGG	5019
1514	UUGUACCG A CCGUCCAC	1997	GTGGACGG GGCTAGCTACAACGA CGGTACAA	5020
1521	GACCGUCC A CGGGGCGC	1998	GCGCCCCG GGCTAGCTACAACGA GGACGGTC	5021
1530	CGGGGCGC A CCUCUCUU	1999	AAGAGAGG GGCTAGCTACAACGA GCGCCCCG	5022
1540	CUCUCUUU A CGCGGACU	2000	AGTCCGCG GGCTAGCTACAACGA AAAGAGAG	5023
1546	UUACGCGG A CUCCCCGU	2001	ACGGGGAG GGCTAGCTACAACGA CCGCGTAA	5024
1567	GCCUUCUC A UCUGCCGG	2002	CCGGCAGA GGCTAGCTACAACGA GAGAAGGC	5025
1576	UCUGCCGG A CCGUGUGC	2003	GCACACGG GGCTAGCTACAACGA CCGGCAGA	5026
1585	CCGUGUGC A CUUCGCUU	2004	AAGCGAAG GGCTAGCTACAACGA GCACACGG	5027
1595	UUCGCUUC A CCUCUGCA	2005	TGCAGAGG GGCTAGCTACAACGA GAAGCGAA	5028
1603	ACCUCUGC A CGUCGAU	2006	ATGCGACG GGCTAGCTACAACGA GCAGAGGT	5029
1610	CACGUCGC A UGGAGACC	2007	GGTCTCCA GGCTAGCTACAACGA GCGACGTG	5030
1616	GCAUGGAG A CCACCGUG	2008	CACGGTGG GGCTAGCTACAACGA CTCCATGC	5031
1619	UGGAGACC A CCGUGAAC	2009	GTTACCGG GGCTAGCTACAACGA GGTCTCCA	5032
1626	CACCGUGA A CCGCCACA	2010	TGTGGGCG GGCTAGCTACAACGA TCACGGTG	5033
1638	CCACAGGA A CCUGCCCA	2011	TGGGCGAG GGCTAGCTACAACGA TCCTGTGG	5034
1656	GGUCUUGC A UAAGAGGA	2012	TCCTCTTA GGCTAGCTACAACGA GCAAGACC	5035
1664	AUAAGAGG A CUCUUGGA	2013	TCCAAGAG GGCTAGCTACAACGA CCTCTTAT	5036
1672	ACUCUUGG A CUUUCAGC	2014	GCTGAAAG GGCTAGCTACAACGA CCAAGAGT	5037
1682	UUUCAGCA A UGUCAACG	2015	CGTTGACA GGCTAGCTACAACGA TGCTGAAA	5038
1688	CAAUGUCA A CGACCGAC	2016	GTCGGTCG GGCTAGCTACAACGA TGACATTG	5039
1691	UGUCAACG A CCGACCUU	2017	AAGGTCGG GGCTAGCTACAACGA CGTTGACA	5040
1695	AACGACCG A CCUUGAGG	2018	CCTCAAGG GGCTAGCTACAACGA CGGTCGTT	5041
1705	CUUGAGGC A UACUUCAA	2019	TTGAAGTA GGCTAGCTACAACGA GCCTCAAG	5042
1707	UGAGGCAU A CUUCAAAAG	2020	CTTTGAAG GGCTAGCTACAACGA ATGCCTCA	5043
1716	CUUCAAAAG A CUGUGUGU	2021	ACACACAG GGCTAGCTACAACGA CTTTGAAG	5044
1728	UGUGUUUA A UGAGUGGG	2022	CCCCTCA GGCTAGCTACAACGA TAAACACA	5045
1774	GUCUUGU A CUAGGAGG	2023	CCTCCTAG GGCTAGCTACAACGA ACAAAGAC	5046
1791	CUGUAGGC A UAAAUUGG	2024	CCAATTTA GGCTAGCTACAACGA GCCTACAG	5047
1795	AGGCAUAA A UUGGUGUG	2025	CACACCAA GGCTAGCTACAACGA TTATGCCT	5048
1807	GUGUGUUC A CCAGCACC	2026	GGTGCTGG GGCTAGCTACAACGA GAACACAC	5049
1813	UCACCAGC A CCAUGCAA	2027	TTGCATGG GGCTAGCTACAACGA GCTGGTGA	5050
1816	CCAGCACC A UGCAACUU	2028	AAGTTGCA GGCTAGCTACAACGA GGTGCTGG	5051
1821	ACCAUGCA A CUUUUUCA	2029	TGAAAAAG GGCTAGCTACAACGA TGCATGGT	5052
1829	ACUUUUUC A CCUCUGCC	2030	GGCAGAGG GGCTAGCTACAACGA GAAAAAGT	5053
1840	UCUGCCUA A UCAUCUCA	2031	TGAGATGA GGCTAGCTACAACGA TAGGCAGA	5054
1843	GCCUAAUC A UCUCUUGU	2032	ACATGAGA GGCTAGCTACAACGA GATTAGGC	5055
1848	AUCAUCUC A UGUUCAUG	2033	CATGAACA GGCTAGCTACAACGA GAGATGAT	5056
1854	UCAUGUUC A UGUCCUAC	2034	GTAGGACA GGCTAGCTACAACGA GAACATGA	5057
1861	CAUGUCCU A CUGUUCAA	2035	TTGAACAG GGCTAGCTACAACGA AGGACATG	5058
1903	UUUGGGGC A UGGACAUU	2036	AATGTCCA GGCTAGCTACAACGA GCCCAAAA	5059
1907	GGGAUGG A CAUUGACC	2037	GGTCAATG GGCTAGCTACAACGA CCATGCCC	5060
1909	GCAUGGAC A UUGACCCG	2038	CGGGTCAA GGCTAGCTACAACGA GTCCATGC	5061

Table 41

1913	GGACAUUG A CCCGUAUA	2039	TATACGGG GGCTAGCTACAACGA CAATGTCC	5062
1919	UGACCCGU A UAAAGAAU	2040	ATTCTTTA GGCTAGCTACAACGA ACGGGTCA	5063
1926	UAUAAAGA A UUUGGAGC	2041	GCTCCAAA GGCTAGCTACAACGA TCTTTATA	5064
1947	GUGGAGUU A CUCUCUUU	2042	AAAGAGAG GGCTAGCTACAACGA AACTCCAC	5065
1967	GCCUUCUG A CUUCUUUC	2043	GAAAGAAG GGCTAGCTACAACGA CAGAAGGC	5066
1981	UUCUUCU A UUCGAGAU	2044	ATCTCGAA GGCTAGCTACAACGA AGAAGGAA	5067
1988	UAUUCGAG A UCUCUCG	2045	CGAGGAGA GGCTAGCTACAACGA CTCGAATA	5068
1997	UCUCUCG A CACCGCCU	2046	AGGCGGTG GGCTAGCTACAACGA CGAGGAGA	5069
1999	UCCUCGAC A CCGCCUCU	2047	AGAGGCGG GGCTAGCTACAACGA GTCGAGGA	5070
2015	UGCUCUGU A UCGGGGGG	2048	CCCCCGGA GGCTAGCTACAACGA ACAGAGCA	5071
2040	UCUCCGGA A CAUUGUUC	2049	GAACAATG GGCTAGCTACAACGA TCCGAGGA	5072
2042	UCCGGAAC A UUGUUCAC	2050	GTGAACAA GGCTAGCTACAACGA GTTCCGGA	5073
2049	CAUUGUUC A CCUCACCA	2051	TGGTGAGG GGCTAGCTACAACGA GAACAATG	5074
2054	UUCACCUC A CCAUACGG	2052	CCGTATGG GGCTAGCTACAACGA GAGGTGAA	5075
2057	ACCUCACC A UACGGCAC	2053	GTGCCGTA GGCTAGCTACAACGA GGTGAGGT	5076
2059	CUCACCAU A CGGCACUC	2054	GAGTGCCG GGCTAGCTACAACGA ATGGTGAG	5077
2064	CAUACGGC A CUCAGGCA	2055	TGCCTGAG GGCTAGCTACAACGA GCCGTATG	5078
2077	GGCAAGCU A UUCUGUGU	2056	ACACAGAA GGCTAGCTACAACGA AGCTTGCC	5079
2098	GUGAGUUG A UGAAUCUA	2057	TAGATTCA GGCTAGCTACAACGA CAACTCAC	5080
2102	GUUGAUGA A UCUAGCCA	2058	TGGCTAGA GGCTAGCTACAACGA TCATCAAC	5081
2110	AUCUAGCC A CCUGGGUG	2059	CACCCAGG GGCTAGCTACAACGA GGCTAGAT	5082
2126	GGGAAGUA A UUUGGAAG	2060	CTTCCAAA GGCTAGCTACAACGA TACTTCCC	5083
2135	UUUGGAAG A UCCAGCAU	2061	ATGTGAAA GGCTAGCTACAACGA CTTCAAA	5084
2142	GAUCCAGC A UCCAGGGA	2062	TCCCTGGA GGCTAGCTACAACGA GCTGGATC	5085
2151	UCCAGGGA A UUAGUAGU	2063	ACTACTAA GGCTAGCTACAACGA TCCTGGA	5086
2165	AGUCAGCU A UGUCAACG	2064	CGTTGACA GGCTAGCTACAACGA AGCTGACT	5087
2171	CUAUGUCA A CGUUAUA	2065	TATTAACG GGCTAGCTACAACGA TGACATAG	5088
2177	CAACGUUA A UAUGGGCC	2066	GGCCCAT A GGCTAGCTACAACGA TAACGTTG	5089
2179	ACGUUAU A UGGGCCUA	2067	TAGGCCCA GGCTAGCTACAACGA ATTAACGT	5090
2191	GCCUAAAA A UCAGACAA	2068	TTGTCTGA GGCTAGCTACAACGA TTTAGGC	5091
2196	AAAUCAG A CAACUAU	2069	AATAGTTG GGCTAGCTACAACGA CTGATTTT	5092
2199	AUCAGACA A CUAUUGUG	2070	CACAATAG GGCTAGCTACAACGA TGTCTGAT	5093
2202	AGACAACU A UUGUGGUU	2071	AACCACAA GGCTAGCTACAACGA AGTTGTCT	5094
2213	GUGGUUUC A CAUUCCU	2072	AGGAAATG GGCTAGCTACAACGA GAAACCAC	5095
2215	GGUUCAC A UUUCUGU	2073	ACAGGAAA GGCTAGCTACAACGA GTGAAACC	5096
2227	CCUGUCU A CUUUUGG	2074	CCCAAAAG GGCTAGCTACAACGA AAGACAGG	5097
2242	GGCAGAA A CUGUUCU	2075	AAGAACAG GGCTAGCTACAACGA TTCTCGCC	5098
2253	GUUCUGA A UAUUUGU	2076	ACCAAATA GGCTAGCTACAACGA TCAAGAAC	5099
2255	UCUGAAU A UUUGGUGU	2077	ACACCAA GGCTAGCTACAACGA ATTCAAGA	5100
2278	GAGUGUGG A UUCGCACU	2078	AGTGCGAA GGCTAGCTACAACGA CCACACTC	5101
2284	GGAUUCG A CUCCUCCU	2079	AGGAGGAG GGCTAGCTACAACGA GCGAATCC	5102
2295	CCUCCUGC A UAUAGACC	2080	GGTCTATA GGCTAGCTACAACGA GCAGGAGG	5103
2297	UCCUGCAU A UAGACCAC	2081	GTGGTCTA GGCTAGCTACAACGA ATGCAGGA	5104
2301	GCAUAUAG A CCACCAA	2082	TTTGGTGG GGCTAGCTACAACGA CTATATGC	5105
2304	UAUAGACC A CCAAUUGC	2083	GCATTTGG GGCTAGCTACAACGA GTTCTATA	5106
2309	ACCACCAA A UGCCCCUA	2084	TAGGGGCA GGCTAGCTACAACGA TTGGTGGT	5107
2317	AUGCCCCU A UCUAUCA	2085	TGATAAGA GGCTAGCTACAACGA AGGGGCAT	5108
2322	CCUAUCU A UCAACACU	2086	AGTGTGTA GGCTAGCTACAACGA AAGATAGG	5109
2326	UCUAUCA A CACUCCG	2087	CGGAAGTG GGCTAGCTACAACGA TGATAAGA	5110
2328	UUAUCAAC A CUUCCGGA	2088	TCCGGAAG GGCTAGCTACAACGA GTTGATAA	5111
2338	UUCCGGAA A CUACUGUU	2089	AACAGTAG GGCTAGCTACAACGA TTCCGGAA	5112

Table 41

2341	CGGAAACU A CUGUUGUU	2090	AACAACAG GGCTAGCTACAACGA AGTTTCCG	5113
2352	GUUGUUG A CGAAGAGG	2091	CCTCTTCG GGCTAGCTACAACGA CTAACAAC	5114
2380	GAAGAAGA A CUCCCUCG	2092	CGAGGGAG GGCTAGCTACAACGA TCTTCTTC	5115
2397	CCUCGCAG A CGAAGGUC	2093	GACCTTCG GGCTAGCTACAACGA CTGCGAGG	5116
2409	AGGUCUCA A UCGCCGCG	2094	CGCGGCGA GGCTAGCTACAACGA TGAGACCT	5117
2427	CGCAGAAG A UCUCAAUC	2095	GATTGAGA GGCTAGCTACAACGA CTTCTGCG	5118
2433	AGAUCUCA A UCUCGGGA	2096	TCCCGAGA GGCTAGCTACAACGA TGAGATCT	5119
2442	UCUCGGGA A UCUCAAUG	2097	CATTGAGA GGCTAGCTACAACGA TCCCGAGA	5120
2448	GAUUCUCA A UGUUAGUA	2098	TACTAACA GGCTAGCTACAACGA TGAGATTC	5121
2456	AUGUUGU A UUCUUGG	2099	CCAAGGAA GGCTAGCTACAACGA ACTAACAT	5122
2465	UUCUUGG A CACAUAA	2100	CTTATGTG GGCTAGCTACAACGA CCAAGGAA	5123
2467	CCUUGGAC A CAUAAGGU	2101	ACCTTATG GGCTAGCTACAACGA GTCCAAGG	5124
2469	UUGGACAC A UAAGGUGG	2102	CCACCTTA GGCTAGCTACAACGA GTGTCCAA	5125
2481	GGUGGGAA A CUUACGG	2103	CCGTAAAG GGCTAGCTACAACGA TTCCCAAC	5126
2486	GAACUUU A CGGGGCUU	2104	AAGCCCCG GGCTAGCTACAACGA AAAGTTTC	5127
2496	GGGGCUU A UUCUUCUA	2105	TAGAAGAA GGCTAGCTACAACGA AAAGCCCC	5128
2504	AUUCUUCU A CGGUACCU	2106	AGGTACCG GGCTAGCTACAACGA AGAAGAAT	5129
2509	UCUACGGU A CCUUGCUU	2107	AAGCAAGG GGCTAGCTACAACGA ACCGTAGA	5130
2520	UUGCUUUA A UCCUAAAU	2108	ATTTAGGA GGCTAGCTACAACGA TAAAGCAA	5131
2527	AAUCCUAA A UGGCAAAC	2109	GTTTGCCA GGCTAGCTACAACGA TTAGGATT	5132
2534	AAUGGCAA A CUCCUUCU	2110	AGAAGGAG GGCTAGCTACAACGA TTGCCATT	5133
2550	UUUUCUG A CAUUCAUU	2111	AATGAATG GGCTAGCTACAACGA CAGGAAAA	5134
2552	UUCUGAC A UUCAUUG	2112	CAAATGAA GGCTAGCTACAACGA GTCAGGAA	5135
2556	UGACAUUC A UUUGCAGG	2113	CCTGCAAA GGCTAGCTACAACGA GAATGTCA	5136
2568	GCAGGAGG A CAUUGUUG	2114	CAACAATG GGCTAGCTACAACGA CCTCCTGC	5137
2570	AGGAGGAC A UUGUUGAU	2115	ATCAACAA GGCTAGCTACAACGA GTCCTCCT	5138
2577	CAUUGUUG A UAGAUGUA	2116	TACATCTA GGCTAGCTACAACGA CAACAATG	5139
2581	GUUGAUAG A UGUAAGCA	2117	TGCTTACA GGCTAGCTACAACGA CTATCAAC	5140
2590	UGUAAGCA A UUUGUGG	2118	CCCACAAA GGCTAGCTACAACGA TGCTTACA	5141
2606	GGCCCUU A CAGUAAAU	2119	ATTTACTG GGCTAGCTACAACGA AAGGGGCC	5142
2613	UACAGUAA A UGAAACA	2120	TGTTTTCA GGCTAGCTACAACGA TTACTGTA	5143
2619	AAAUAAA A CAGGAGAC	2121	GTCTCCTG GGCTAGCTACAACGA TTTCATTT	5144
2626	AACAGGAG A CUUAAAUU	2122	AATTTAAG GGCTAGCTACAACGA CTCCTGTT	5145
2632	AGACUUA A UUAACUUA	2123	ATAGTTAA GGCTAGCTACAACGA TTAAGTCT	5146
2636	UUAAAUUA A CUAUGCCU	2124	AGGCATAG GGCTAGCTACAACGA TAATTTAA	5147
2639	AAUUAACU A UGCCUGCU	2125	AGCAGGCA GGCTAGCTACAACGA AGTTAATT	5148
2655	UAGGUUUU A UCCCAAUG	2126	CATTGGGA GGCTAGCTACAACGA AAAACCTA	5149
2661	UUAUCCCA A UGUUACUA	2127	TAGTAACA GGCTAGCTACAACGA TGGGATAA	5150
2666	CAAUGUU A CUAAAUUA	2128	ATATTTAG GGCTAGCTACAACGA AACATTGG	5151
2671	GUUACUAA A UAUUUGCC	2129	GGCAAATA GGCTAGCTACAACGA TTAGTAAC	5152
2673	UACUAAAU A UUUGCCCU	2130	AGGGCAAA GGCTAGCTACAACGA ATTTAGTA	5153
2685	GCCCUUAG A UAAAGGGA	2131	TCCCTTTA GGCTAGCTACAACGA CTAAGGGC	5154
2693	AUAAAGGG A UCAAACCG	2132	CGGTTTGA GGCTAGCTACAACGA CCCTTTAT	5155
2698	GGGAUCAA A CCGUAUUA	2133	TAATACGG GGCTAGCTACAACGA TTGATCCC	5156
2703	CAAACCGU A UUAUCCAG	2134	CTGGATAA GGCTAGCTACAACGA ACGGTTTG	5157
2706	ACCGUAUU A UCCAGAGU	2135	ACTCTGGA GGCTAGCTACAACGA AATACGGT	5158
2715	UCCAGAGU A UGUAGUUA	2136	TAATAACA GGCTAGCTACAACGA ACTCTGGA	5159
2724	UGUAGUUA A UCAUUAU	2137	AGTAATGA GGCTAGCTACAACGA TAACTACA	5160
2727	AGUUAUUC A UUAUUUCC	2138	GGAAGTAA GGCTAGCTACAACGA GATTAACT	5161
2730	UAAUCAUU A CUUCCAGA	2139	TCTGGAAG GGCTAGCTACAACGA AATGATTA	5162
2738	ACUUCAG A CGCGACAU	2140	ATGTCGCG GGCTAGCTACAACGA CTGGAAGT	5163

Table 4 I

2743	CAGACGCG A CAUUAUUU	2141	AAATAATG GGCTAGCTACAACGA CGCGTCTG	5164
2745	GACGCGAC A UUAUUUAC	2142	GTAAATAA GGCTAGCTACAACGA GTCGCGTC	5165
2748	GCGACAUU A UUUACACA	2143	TGTGTAAA GGCTAGCTACAACGA AATGTCGC	5166
2752	CAUUAUUU A CACACUCU	2144	AGAGTGTG GGCTAGCTACAACGA AAATAATG	5167
2754	UUAUUUAC A CACUCUUU	2145	AAAGAGTG GGCTAGCTACAACGA GTAAATAA	5168
2756	AUUUACAC A CUCUUUGG	2146	CCAAAGAG GGCTAGCTACAACGA GTGTAAAT	5169
2774	AGGCGGGG A UCUUAUUAU	2147	ATATAAGA GGCTAGCTACAACGA CCCC GCCT	5170
2779	GGGAUCUU A UAUAAAAG	2148	CTTTTATA GGCTAGCTACAACGA AAGATCCC	5171
2781	GAUCUUUAU A UAAAAGAG	2149	CTCTTTTA GGCTAGCTACAACGA ATAAGATC	5172
2795	GAGAGUCC A CACGUAGC	2150	GCTACGTG GGCTAGCTACAACGA GGA CTCTC	5173
2797	GAGUCCAC A CGUAGCGC	2151	GCGCTACG GGCTAGCTACAACGA GTGGACTC	5174
2809	AGCGCCUC A UUUUGCGG	2152	CCGCAAAA GGCTAGCTACAACGA GAGGCGCT	5175
2821	UGCGGGUC A CCAUAUUC	2153	GAATATGG GGCTAGCTACAACGA GACCCGCA	5176
2824	GGGUCACC A UAUUCUUG	2154	CAAGAATA GGCTAGCTACAACGA GGTGACCC	5177
2826	GUCACCAU A UUCUUGGG	2155	CCCAAGAA GGCTAGCTACAACGA ATGGTGAC	5178
2836	UCUUGGGA A CAAGAUCU	2156	AGATCTTG GGCTAGCTACAACGA TCCCAAGA	5179
2841	GGAACAAG A UCUACAGC	2157	GCTGTAGA GGCTAGCTACAACGA CTGTTC	5180
2845	CAAGAUCU A CAGCAUGG	2158	CCATGCTG GGCTAGCTACAACGA AGATCTTG	5181
2850	UCUACAGC A UGGGAGGU	2159	ACCTCCCA GGCTAGCTACAACGA GCTGTAGA	5182
2870	UCUCCAA A CCUCGAAA	2160	TTTCGAGG GGCTAGCTACAACGA TTGGAAGA	5183
2883	GAAAAGGC A UGGGACAA	2161	TGTCCCAA GGCTAGCTACAACGA GCCTTTTC	5184
2889	GCAUGGGG A CAAAUUCU	2162	AAGATTTG GGCTAGCTACAACGA CCCCATGC	5185
2893	GGGACAA A UCUUUCUG	2163	CAGAAAGA GGCTAGCTACAACGA TTGTCCTC	5186
2908	UGUCCCCA A UCCCCUGG	2164	CCAGGGGA GGCTAGCTACAACGA TGGGACAA	5187
2918	CCCCUGGG A UUCUCCCC	2165	GGGAAGAA GGCTAGCTACAACGA CCCAGGGG	5188
2929	CUUCCCCG A UCAUCAGU	2166	ACTGATGA GGCTAGCTACAACGA CGGGGAAG	5189
2932	CCCCGAUC A UCAGUUGG	2167	CCAAGTGA GGCTAGCTACAACGA GATCGGGG	5190
2941	UCAGUUGG A CCCUGCAU	2168	ATGCAGGG GGCTAGCTACAACGA CCAAGTGA	5191
2948	GACCCUGC A UUCAAGC	2169	GCTTTGAA GGCTAGCTACAACGA GCAGGGTC	5192
2959	CAAAGCCA A CUCAGUAA	2170	TTACTGAG GGCTAGCTACAACGA TGGCTTTG	5193
2968	CUCAGUAA A UCCAGAUU	2171	AATCTGGA GGCTAGCTACAACGA TTACTGAG	5194
2974	AAAUCCAG A UUGGGACC	2172	GGTCCCAA GGCTAGCTACAACGA CTGGATTT	5195
2980	AGAUUGGG A CCUCAACC	2173	GGTTGAGG GGCTAGCTACAACGA CCAATCT	5196
2986	GGACCUCA A CCCGCACA	2174	TGTGCGGG GGCTAGCTACAACGA TGAGGTCC	5197
2998	GCACAAGG A CAACUGGC	2175	GCCAGTTG GGCTAGCTACAACGA CCTGTGTC	5198
3001	CAAGGACA A CUGGCCGG	2176	CCGGCCAG GGCTAGCTACAACGA TGTCTTTG	5199
3010	CUGGCCGG A CGCCAACA	2177	TGTTGGCG GGCTAGCTACAACGA CCGGCCAG	5200
3016	GGACGCCA A CAAGGUGG	2178	CCACCTTG GGCTAGCTACAACGA TGGCGTCC	5201
3035	GUGGGAGC A UUCGGGCC	2179	GGCCCCGA GGCTAGCTACAACGA GCTCCCAC	5202
3051	CAGGGUUC A CCCUCCCC	2180	GGGAGGGG GGCTAGCTACAACGA GAACCTTG	5203
3061	CCCUCCCC A UGGGGGAC	2181	GTCCCCCA GGCTAGCTACAACGA GGGGAGGG	5204
3068	CAUGGGGG A CUGUUGGG	2182	CCCAACAG GGCTAGCTACAACGA CCCCATG	5205
3088	GAGCCUC A CGCUCAGG	2183	CCTGAGCG GGCTAGCTACAACGA GAGGGCTC	5206
3101	CAGGGCCU A CUCACAAC	2184	GTTGTGAG GGCTAGCTACAACGA AGGCCCTG	5207
3105	GCCUACUC A CAACUGUG	2185	CACAGTTG GGCTAGCTACAACGA GAGTAGGC	5208
3108	UACUCACA A CUGUGCCA	2186	TGGCACAG GGCTAGCTACAACGA TGTGAGTA	5209
3138	CUGCCUCC A CCAAUCCG	2187	CCGATTGG GGCTAGCTACAACGA GGAGGCAG	5210
3142	CUCCACCA A UCGGACAGU	2188	ACTGCCGA GGCTAGCTACAACGA TGGTGGAG	5211
3165	GGCAGCCU A CUCCCUUA	2189	TAAGGGAG GGCTAGCTACAACGA AGGCTGCC	5212
3173	ACUCCCUU A UCUCACC	2190	GGTGGAGA GGCTAGCTACAACGA AAGGGAGT	5213
3179	UUAUCUCC A CCUCUAAG	2191	CTTAGAGG GGCTAGCTACAACGA GGAGATAA	5214

Table 4 i

3190	UCUAAGGG A CACUCAUC	2192	GATGAGTG GGCTAGCTACAACGA CCCTTAGA	5215
3192	UAAGGGAC A CUCAUCCU	2193	AGGATGAG GGCTAGCTACAACGA GTCCCTTA	5216
3196	GGACACUC A UCCUCAGG	2194	CCTGAGGA GGCTAGCTACAACGA GAGTGTCC	5217
3207	CUCAGGCC A UGCAGUGG	2195	CCACTGCA GGCTAGCTACAACGA GGCCTGAG	5218

Input Sequence = AF100308. Cut Site = YG/M or UG/U.

Stem Length = 8 . Core Sequence = GGCTAGCTACAACGA

AF100308 (Hepatitis B virus strain 2-18, 3215 bp)

Table 42

Table 42: Human HBV Amberzyme Ribozyme and Substrate Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
61	ACUUUCU G CUGGUGG	1448	GCCACCAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AGGAAAGU	5219
87	GGAACAGU G AGCCUUG	1449	GCAGGCU GGAGGAAACUCC CU UCAAGGACAUUGUCCGG ACUGUUC	5220
94	UGAGCCCU G CUCAGAAU	1450	AUCUGAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AGGCUCA	5221
112	CUGUCUCU G CCAUAUCG	1451	CGAUAUGG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AGAGACAG	5222
132	AUCUUAUC G AAGACUGG	1452	CCAGUCUU GGAGGAAACUCC CU UCAAGGACAUUGUCCGG GAUAAGAU	5223
153	CCUGUACC G AACUGGA	1453	UCCAUGUU GGAGGAAACUCC CU UCAAGGACAUUGUCCGG GGUACAGG	5224
169	AGAACAU G CAUCAGGA	1454	UCCUGAUG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG GAUGUUCU	5225
192	GGACCCCU G CUCGUGUU	1455	AACACGAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AGGGGUCC	5226
222	UUCUUGUU G AAAAAAU	1456	AUUUUUGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AACAAAGAA	5227
315	CAAAAUUC G CAGUCCCA	1457	UGGGACUG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG GAAUUUUG	5228
374	UGGUUAUC G CUGGAUGU	1458	ACAUCAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG GAUAACCA	5229
387	AUGUGUCU G CGCGUUG	1459	AAACGCG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AGACACAU	5230
410	CUUCUCU G CAUCCUGC	1460	GCAGGUG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AGAGGAAG	5231
417	UGCAUCCU G CUGCUAUG	1461	CAUAGCAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AGGAUGCA	5232
420	AUCCUGCU G CUAUGCCU	1462	AGGCAUAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AGCAGGAU	5233
425	GCUGCUAU G CCUCAUGU	1463	AGAUGAGG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AUAGCAGC	5234
468	GGUAUGUU G CCGGUTUG	1464	CAAAACGG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AACAUACC	5235
518	CGGACCAU G CAAAAACU	1465	AGGUUUUG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AUGGUCCG	5236
527	CAAAACCU G CACAAACU	1466	GAGUUUG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AGGUUUUG	5237
538	CAACUCCU G CUCAAAGGA	1467	UCCUUGAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AGGAGUUG	5238
569	CUCAUGUU G CUGUACAA	1468	UUGUACAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AACAUAG	5239
596	CGGAAACU G CACCUGUA	1469	UACAGGUG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AGUUUCCG	5240
631	GGGCUUUC G CAAAAUAC	1470	GUUUUUUG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG GAAAGCCC	5241
687	UUACUAGU G CCAUUUGU	1471	ACAAAUGG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG ACTAGUAA	5242
747	AUAUGGAU G AUGUGGUU	1472	AACACAU GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AUCCAUAU	5243
783	AACAUCUU G AGUCCUUU	1473	AAGGACU GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AAGAUGUU	5244
795	CCUUUAU G CCGUUGUU	1474	AACAGCGG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AUAAGGG	5245
798	UUUAUGCC G CUGUUUAC	1475	GGUAACAG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG GGCAUAAA	5246
911	GGCAUUAU G CCACAGGA	1476	UCCUGUGG GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AAUGUCC	5247
978	GGCCUAUU G AUUGGAAA	1477	UUUCCAAU GGAGGAAACUCC CU UCAAGGACAUUGUCCGG AAUAGGCC	5248

Table 42

997	AUGUCAAC G AAUUGUGG	1478	CCACAAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUUGACAU	5249
1020	UGGGGUU G CGGCCCU	1479	AGGGCGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AAACCCCA	5250
1023	GGUUGCC G CCCUUUC	1480	GAAAGGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGCACACC	5251
1034	CCUUUAC G CAAUGUGG	1481	CCACAUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUGAAAGG	5252
1050	GAUAUCU G CUUUAUUG	1482	CAUUAAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGAUUAUC	5253
1058	GCUUUAAU G CCUUUAUA	1483	UAUUAAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUUAAAGC	5254
1068	CUUUAU G CAUGCAUA	1484	UAUGCAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUUAAAG	5255
1072	AUAUGAU G CAUACAAG	1485	CUUGUAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUGCAUAU	5256
1103	ACUUUCU G CCACTUUA	1486	UAAGUUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GAGAAAGU	5257
1139	CAGUAUG G AACCUUUA	1487	UAAAGGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG ACAUACUG	5258
1155	ACCCCGU G CUCGGCAA	1488	UUGCCGAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AACGGGUG	5259
1177	UGGUCUAU G CCAAGUGU	1489	ACACUUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUAGACCA	5260
1188	AAGUGUUU G CUGACGCA	1490	UGCUCAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AAACACUU	5261
1191	UGUUUGU G ACGCAACC	1491	GGUUGCU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGCAAACA	5262
1194	UUGCUGAC G CAACCCCC	1492	GGGGUUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUCAGCAA	5263
1234	CCAUCAG G CAUGCGUG	1493	CACGAUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GCUGAUGG	5264
1238	CAGCGAU G CGUGGAAC	1494	GUUCCAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUGCGCUG	5265
1262	UCUCCUCU G CCGAUCCA	1495	UGGAUCG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGAGGAGA	5266
1265	CCUCUGCC G AUCCAUAU	1496	GUUUGAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGCAGAGG	5267
1275	UCCAUAU G CGGAACUC	1497	GAGUCCG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGUUUGGA	5268
1290	UCCUAGCC G CUUGUUAU	1498	AAACAAAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGCUAGGA	5269
1299	CUUGUUU G CUCGCAGC	1499	GCUGCGAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AAAACAAG	5270
1303	UUUUGCUC G CAGCAGGU	1500	ACCUGCUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GAGCAAAA	5271
1335	UCGGGACU G ACAAUUCU	1501	AGAUUUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGUCCCGA	5272
1349	UCUGUCGU G CUCUCCCG	1502	CGGAGAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG ACGACAGA	5273
1357	GCUCUCC G CAAUAUA	1503	UAUAUUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGGAGAGC	5274
1382	CCAUGGCU G CUAGGCUU	1504	CAGCCUAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGCCAUUG	5275
1392	UAGGCGU G CUGCCAAC	1505	GUUGCAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG ACAGCCUA	5276
1395	GCUGUGU G CCNACTUG	1506	CCAGUUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGCACAGC	5277
1411	GAUCCUAC G CGGACGU	1507	ACGUCCG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUAGGAUC	5278
1442	CCGUCCG G CUGAAUCC	1508	GAUUCAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GCCGACGG	5279
1445	UCGGCGU G AUUCCGCG	1509	GCGGUAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGCGCCGA	5280
1452	UGAAUCC G CGGACGAC	1510	GUCGUCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGGAUUCA	5281
1458	CCGGGAC G ACCCTUCC	1511	GGAGGGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUCCCGGG	5282

Table 42

1474	CCGGGGCC G CUUGGGG	1512	GCCCCAAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GGCCCCGG	5283
1489	GCUCUACC G CCGCUUC	1513	GAAGCGGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GGUAGAGC	5284
1493	UACCGCCC G CUUCUCG	1514	CGGAGAAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GGGCGGUA	5285
1501	GCUCUCC G CCUAUUGU	1515	ACAAUAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GGAGAAGC	5286
1513	AUUGUACC G ACCGUCCA	1516	UGGACGGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GGUACAAU	5287
1528	CACGGGGC G CACCUUC	1517	GAGAGGUG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GCCCCGUG	5288
1542	CUUUUAC G CGGACUCC	1518	GGAGUCCG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GUAAAGAG	5289
1559	CCGUCUGU G CCUUCUCA	1519	UGAGAAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG ACAGACGG	5290
1571	UCUCAUCU G CCGGACCG	1520	CGGUCCGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGAUGAGA	5291
1583	GACCGUGU G CACUUCGC	1521	GCGAAGUG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG ACACGGUC	5292
1590	UGCACUUC G CUUACCU	1522	AGGUGAAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GAAGUGCA	5293
1601	UCACCUUC G CACGUCGC	1523	GCGACGUG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGAGGUGA	5294
1608	UGCACGUC G CAUGGAGA	1524	UCUCCAUG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GACGUGCA	5295
1624	ACCACCGU G AACGCCCA	1525	UGGGCGUU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG ACGGUGGU	5296
1628	CCGUGAAC G CCCACAGG	1526	CCUGUGGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GUTUCACGG	5297
1642	AGGAACCU G CCAAGGU	1527	ACCUUGGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGGUUCCU	5298
1654	AAGGUCUU G CAUAAGAG	1528	CUCUUAUG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AAGACCUU	5299
1690	AUGUCAAC G ACCGACCU	1529	AGGUCGGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GUUGACAU	5300
1694	CAACGACC G ACCUUGAG	1530	CUCAAGGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GGUCGUUG	5301
1700	CCGACCUU G AGGCAUAC	1531	GU AUGCCU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AAGGUCGG	5302
1730	UGUUAAU G AGUGGGAG	1532	CUCCCAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUUAACA	5303
1818	AGCACCAU G CAACUUUU	1533	AAAAGUUG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUGGUGCU	5304
1835	UCACCUUC G CCUAUACA	1534	UGAUUAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGAGGUGA	5305
1883	CAAGCUGU G CCUUGGGU	1535	ACCCAAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG ACAGCUUG	5306
1912	UGGACAUU G ACCCGUAU	1536	AUACGGGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AAUGUCCA	5307
1959	UCUUUUU G CCUUCUGA	1537	UCAGAAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AAAAAGA	5308
1966	UGCCUUUC G ACUUCUUU	1538	AAAGAAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGAAGGCA	5309
1985	UUCUAUUC G AGAUUCUC	1539	GGAGAUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GAAUAGAA	5310
1996	AUCUCCUC G ACACCGCC	1540	GGCGGUGU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GAGGAGAU	5311
2002	UCGACACC G CCUCUGCU	1541	AGCAGAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG GGUGUGGA	5312
2008	CCGCCUCU G CUCUGUAU	1542	AUACAGAG GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AGAGGCGG	5313
2092	GUUGGGGU G AGUUGAUG	1543	CAUCARAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG ACCCCAAC	5314
2097	GGUGAGUU G AUGAUAUCU	1544	AGAUCAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AACUCACC	5315
2100	GAGUUGAU G AAUCUAGC	1545	GCUAUAU GGAGGAAACUCC CU UCAAGGACAUCGUCCCGG AUCACACU	5316

Table 42

2237	UUUUGGC G AGAAACUG	1546	CAGUUUCU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GCCCAAAA	5317
2251	CUGUUCUU G AAUAUUUG	1547	CAAAUAUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGAACACAG	5318
2282	GUGGAUUC G CACUCCUC	1548	GAGGAGUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GAUCCAC	5319
2293	CUCCUCCU G CAUAUAGA	1549	UCUAUAUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGGAGGAG	5320
2311	CACCAAAU G CCCUAUC	1550	GAUAGGGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUTUUGGUG	5321
2354	UGUAAGAC G AGAGGGCA	1551	UGCCUCUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUCUAACA	5322
2388	ACUCCUUC G CTUCGCAG	1552	CUGCGAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GAGGAGU	5323
2393	CUCGCCUC G CAGACGAA	1553	UUCGUUCG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GAGGCGAG	5324
2399	UCGCAGAC G AAGGUCUC	1554	GAGACCUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUCUGCGA	5325
2412	UCUCAAU G CCGCGUCG	1555	CGACGGGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GAUUGAGA	5326
2415	CAAUCCGC G CGUCGCAG	1556	CUGCGACG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GCGCAUUG	5327
2420	GCCGCGUC G CAGAAGAU	1557	AUCUUCUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GACGCGGC	5328
2514	GGUACCUU G CUUUAUUC	1558	GAUUAAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AAGGUACC	5329
2549	CUUUUCCU G ACAUUCAU	1559	AUGAAUGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGGAAAAG	5330
2560	AUUCAUUU G CAGGAGGA	1560	UCCUCCUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AAUUGAAU	5331
2576	ACAUUGUU G AUAGAUGU	1561	ACAUCUAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AACAAUGU	5332
2615	CAGUAAAU G AAAACAGG	1562	CCUGUUUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUUUACUG	5333
2641	UUAACUAU G CCUGCUAG	1563	CUAGCAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUAGUUA	5334
2645	CUAUGCCU G CUAGGUUU	1564	AAACCUAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGGCAUAG	5335
2677	AAUAUUU G CCUUAAGA	1565	UCUAAGGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AAUAUUUU	5336
2740	UCCAGAC G CGACAUUA	1566	UAUUGUCG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUCUGGAA	5337
2742	CCAGACGC G ACAUUAUU	1567	AAUAUUGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GCGUCUGG	5338
2804	CACGUAGC G CCUCAUUU	1568	AAUAGAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GTUACGUG	5339
2814	CUCAUUUU G CGGGUCAC	1569	GUGACCCG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AAAAUGAG	5340
2875	CAAAACUC G AAAAGGCA	1570	UGCCUUUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GAGGUUUG	5341
2928	UCUUCCCC G AUCAUCAG	1571	CUGAUGAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGGGAAGA	5342
2946	UGGACCCU G CAUUCAAA	1572	UUUGAUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGGUUCCA	5343
2990	CUCAACCC G CACAAGGA	1573	UCCUUUGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGUUUGAG	5344
3012	GGCCGGAC G CCAACAAG	1574	CUUGUUUG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUCCGGCC	5345
3090	GCCCUAC G CUCAGGC	1575	CCCCUGAG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUGAGGGC	5346
3113	ACAACUUG G CCAGCAGC	1576	GTUCUGGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG ACAGUUGU	5347
3132	CUCCUCCU G CCUCCACC	1577	GGUGAGG GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGGAGGAG	5348
51	AGGGCCCU G UACUJUCC	1578	GGAAAGUA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGGGCCCU	5349
106	AGAAUACU G UCUCUGCC	1579	GGCAGAGA GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGUAUUCU	5350

Table 42

148	GGGACCCU G UACGGAAC	1580	GUUCGGUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AGGUCUCC	5351
198	CUGCUCGU G UUAACAGGC	1581	GCCUGUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG ACAGCAG	5352
219	UUUUUCUU G UUGACAAA	1582	UUUGUCAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAGAAAA	5353
297	ACACCCGU G UGUUUUGG	1583	CCAAGACA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG ACGGUGU	5354
299	ACCCGUGU G UCUUUGGC	1584	GGCCAAGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG ACACGGGU	5355
347	ACCAACCU G UUGUCCUC	1585	GAGGACAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AGGUUGU	5356
350	AACCUUUU G UCCUCCAA	1586	UUGGAGGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AACAGGU	5357
362	UCCAAUUU G UCCUUGGU	1587	AACCAGGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAUUGGA	5358
381	CGCUGGAU G UGUUCGCG	1588	CGCAGACA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUCCAGCG	5359
383	CUGGAUGU G UCUGCGGC	1589	GCCGCAGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG ACAUCCAG	5360
438	AUCUUCUU G UUGGUUCU	1590	AGAACCAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAGAAGAU	5361
465	CAAGGUUU G UUGCCCGU	1591	ACGGGCAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUACUUG	5362
476	GCCCGUUU G UCCUCUAA	1592	UUAGAGGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAACGGGC	5363
555	ACCUCUAU G UTUCCUCC	1593	GAGGGAAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUAGAGGU	5364
566	UCCUCUAU G UUGCUGUA	1594	UACAGCAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUGAGGGA	5365
572	AUGUUGCU G UACAAAAC	1595	GUUUUGUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AGCAACAU	5366
602	CUGCACCU G UAUUCCCA	1596	UGGGAUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AGGUGCAG	5367
694	UGCCAUUU G UUCAGUGG	1597	CCACUGAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAUUGGCA	5368
724	CCCCACU G UCUGGCUU	1598	AAGCCAGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AGUGGGGG	5369
750	UGGAUGAU G UGGUUUUG	1599	CAAAACCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUCAUCCA	5370
771	CCAAGUCU G UACAACAU	1600	AUUUGUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AGACUUGG	5371
801	AUGCCGCU G UUAACCAU	1601	AUUGGUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AGCGGCAU	5372
818	UUUCUUUU G UCUUUGGG	1602	CCCAAGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAAAGAAA	5373
888	UGGGAUAU G UAAUUGGG	1603	CCCAUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUAUCCCA	5374
927	AACAUAUU G UACAAAAA	1604	UUUUUGUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAUAUGUU	5375
944	AUCAAUUU G UGUUUUAG	1605	CUAAACA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUUUUGAU	5376
946	CAAAAUUU G UUUUAGGA	1606	UCCUAAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG ACAUUUUG	5377
963	AACUUCUU G UAAACAGG	1607	CCUUUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AGGAAGUU	5378
991	GAAGUAUU G UCAACGAA	1608	UUCGUUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUACUUUC	5379
1002	AACGAUUU G UGGGUUUU	1609	AAGACCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAUUCGUU	5380
1039	CACGCAUU G UGGAUUUU	1610	AAUAUCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUUUGCGU	5381
1137	AACAGUAU G UGAACCUU	1611	AAGGUUCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUACUGUU	5382
1184	UGCCAAGU G UTUUGUGA	1612	UCAGCAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG ACUUGGCA	5383
1251	GAACUUUU G UGUUCUCCU	1613	AGGAGACA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAAGGUUC	5384

Table 42

1253	ACUUUUU G UCUCUCU	1614	AGAGGAGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGAAGGU	5385
1294	AGCCGCUU G UUUUGUC	1615	GAGCAAAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGCGGCU	5386
1344	ACAUTUCU G UGUGUCU	1616	GAGCAGGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGAAUUGU	5387
1390	GCUAGGCU G UGUGCCA	1617	UGGAGGCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGCCUAGC	5388
1425	CGUCCUUU G UUUACGUC	1618	GACGUAAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AAAGGACG	5389
1508	CGCCUAUU G UACCGACC	1619	GGUCGGUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AAUAGGCG	5390
1557	CCCCGUCU G UGCCUUCU	1620	AGAAGGCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGACGGGG	5391
1581	CGGACCGU G UGCACUUC	1621	GAAGUGCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG ACGUCCCG	5392
1684	UCAGCAAU G UCAACGAC	1622	GUCGUUGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AUUGCUGA	5393
1719	CAAAGACU G UGUGUUUA	1623	UAAACACA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGUCUUUG	5394
1721	AAGACUGU G UGUUUAAU	1624	AUUAACA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG ACAGUCUU	5395
1723	GACUGUGU G UUUUAUGA	1625	UCAUUAAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG ACACAGUC	5396
1772	AGGUCUUU G UACUAGGA	1626	UCCUAGUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AAAGACCU	5397
1785	AGGAGGCU G UAGGCAUA	1627	UAUGCCUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGCCUCCU	5398
1801	AAAUTUGU G UGUUCACC	1628	GGUGAACA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG ACCAAUUT	5399
1803	AUTGGUGU G UUCACCAG	1629	CUGGUGAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG ACACCAAU	5400
1850	CAUCUCAU G UUCAUGUC	1630	GACAUGAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AUGAGAUG	5401
1856	AUGUJCAU G UCCUACUG	1631	CAGUAGGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AUGAACAU	5402
1864	GUCCUACU G UUCAAGCC	1632	GGCUUGAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGUAGGAC	5403
1881	UCCAAGCU G UGCCUUGG	1633	CCAAGGCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGCUUGGA	5404
1939	GAGCUTUCU G UGGAGUUA	1634	UAACUCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGAAGCUC	5405
2013	UCUGUCUCU G UAUCGGGG	1635	CCCCGAUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGAGCAGA	5406
2045	GGAACAUU G UUCACCCUC	1636	GAGGUGAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AAUGUUCU	5407
2082	GCUAUTUCU G UGUUGGGG	1637	CCCCAACA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGAAUAGC	5408
2084	UAUTUCUGU G UUGGGGUG	1638	CACCCCAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG ACAGAAUA	5409
2167	UCAGCUAU G UCAACGCU	1639	AACGUUGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AUAGCUGA	5410
2205	CAACUAUU G UGGUUUCA	1640	UGAAACCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AAUAGUUG	5411
2222	CAUUTUCU G UCUAUCUU	1641	AAGUAAGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGGAAUUG	5412
2245	GAGAAACU G UUCUUGAA	1642	UUCRAGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGUUUCUC	5413
2262	UAUUGGUGU G UCUUUUGG	1643	CCAAAGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG ACCAAUA	5414
2274	UUUGGAGU G UGGAUUCG	1644	CGAAUCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG ACUCCAAA	5415
2344	AAACUACU G UUGUUAGA	1645	UCUAACA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AGUAGUUU	5416
2347	CUACUGUU G UUAGACGA	1646	UCGUCUAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AACAGUAG	5417
2450	AUCUCAAU G UUAGUAUU	1647	AAUACUAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG AUUGAGAU	5418

Table 42

2573	AGGACAUU G UUAUAGA	1648	UCUAUCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAUUGCUU	5419
2583	UGAUAGAU G UAGCAAU	1649	AUUGCUUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUCUAUCA	5420
2594	AGCAUUU G UGGGGCCC	1650	GGGCCCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAUUGCUU	5421
2663	AUCCCAU G UUAUAAA	1651	UUUAGUAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUUGGGAU	5422
2717	CAGAGUUA G UAGUUAAU	1652	AUUAAUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUACUCUG	5423
2901	AUCUUUCU G UCCCAAU	1653	AUUGGGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGAAAGAU	5424
3071	GGGGGACU G UUGGGGUG	1654	CACCCCAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGUCCCCC	5425
3111	UCACAACU G UGCAGCA	1655	UGCUGGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGUUGUGA	5426
40	AUCCCA G UCAGGGCC	1656	GGCCUGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCUUGGAU	5427
46	GAGUCAG G CCCUGUAC	1657	GUACAGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGACUC	5428
65	UCCUGCUG G UGGCUCCA	1658	UGGAGCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGCAGGA	5429
68	UGCUGGUG G CUCCAGU	1659	AACUGGAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CACCAGCA	5430
74	UGGCUCCA G UUCAGGAA	1660	UUCCUGAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGGAGCCA	5431
85	CAGGAACA G UGAGCCCU	1661	AGGGCUCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGUUCCUG	5432
89	AACAGUGA G CCCUGCUC	1662	GAGCAGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCACUGUU	5433
120	GCCAUAC G UCAAUUU	1663	AAGAUUGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GAUAUGGC	5434
196	CCUGCUC G UGUUACAG	1664	CUGUAACA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GAGCAGGG	5435
205	UGUACAG G CGGGGUU	1665	AAACCCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGUAACA	5436
210	CAGGCGG G UUUUUCU	1666	AAGAAAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCGCCUG	5437
248	ACCACAGA G UCUGAGU	1667	AGUCUAGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCUGUGGU	5438
258	CUAGACUC G UGGUGGAC	1668	GUCCACCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GAGUCUAG	5439
261	GACUCGUG G UGGACUUC	1669	GAAGUCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CACGAGUC	5440
295	GAACACC G UGUGUCUU	1670	AAGACACA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGGUGUUC	5441
305	GUGUCUUG G CCAAAUU	1671	AAUUUGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAAGACAC	5442
318	AUUCGCA G UCCCAAU	1672	AUUGGGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCGAAUU	5443
332	AAUCUCA G UCACUCAC	1673	GUGAGUGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGGAGAUU	5444
368	UUGUCCUG G UUAUCGCU	1674	AGCGUAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGACAA	5445
390	UGUCGGG G GGUUUUU	1675	AUAAACG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGCAGACA	5446
392	UCUGGGC G UUUUAUCA	1676	UGAUAANA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GCCGCAGA	5447
442	UCUUGUG G UUCUUCUG	1677	CAGAANA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAACAAGA	5448
461	CUAUCAG G UAUGUUG	1678	GCAACUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGAUAG	5449
472	UUGUGCCC G UUGUCCU	1679	AGGACAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGGCAACA	5450
506	AACAACA G CACCGGAC	1680	GUCCGUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGGUUGUU	5451
625	CAUCUUG G CUUUGCA	1681	UGCGAAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCAAGAUG	5452

Table 42

648	CUAUGGGA G UGGGCCUC	1682	GAGGCCCA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UCCCAUAG	5453
652	GGGAGUGG G CCUCAGUC	1683	GACUGAGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCACUCCC	5454
658	GGGCCUCA G UCCGUUUC	1684	GAAACGGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGAGGCCC	5455
662	CUCAGUCC G UUUUCUUC	1685	AAGAGAAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG GGACUGAG	5456
672	UUUCUUG G CUCAGUUU	1686	AAACUGAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CAAGAGAA	5457
677	UUGGCUCA G UUUACUAG	1687	CUAGUAAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGAGCCAA	5458
685	GUUUACUA G UGCCAUUU	1688	AAAUCCGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UAGUAAAC	5459
699	UUUGUJCA G UGGUUCGU	1689	ACGAACCA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGAACAAA	5460
702	GUUCAGUG G UUCGUAGG	1690	CCUACGAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CACUGAAC	5461
706	AGUGGUUC G UAGGCUUU	1691	AAGCCCUA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG GAACACAU	5462
711	UUCGUAGG G CUUUCCCC	1692	GGGAAAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCUACGAA	5463
729	ACUGUCUG G CUUUCAGU	1693	ACTGAAAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CAGACAGU	5464
736	GGCUUUCA G UUAUAUGG	1694	CCAUUAAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGAAAGCC	5465
753	AUGAUGUG G UUUUGGGG	1695	CCCCAAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CACAUCAU	5466
762	UUUUGGG G CCAAGUCU	1696	AGACUUGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCCCAAAA	5467
767	GGGGCCAA G UCUGUACA	1697	UGUACAGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUGGCCCC	5468
785	CAUCUUGA G UCCCUUUA	1698	UAAAGGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UCAAGAUG	5469
826	GUCUUUGG G UAUACAUU	1699	AAUGUAUA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCAAAGAC	5470
898	AAUUGGGA G UUGGGGCA	1700	UGCCCCAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UCCCAUUC	5471
904	GAGUUGGG G CACAUUGC	1701	GCAUUGUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCCAACUC	5472
971	GUAAACAG G CUAUUGA	1702	UCAUUGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUGUUUAC	5473
987	AUUGGAAA G UAUUCUAA	1703	UUGACAUU GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUUCCAAU	5474
1006	AAUUGUGG G UCUUUUGG	1704	CCAAAGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCACAAUU	5475
1016	CUUUUGGG G UUUCCCGC	1705	GCGGCAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCCAAAAG	5476
1080	GCAUACAA G CAAAACAG	1706	CUGUUUUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUGUAUGC	5477
1089	CAAAACAG G CUUUUACU	1707	AGUAAAAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUGUUUUU	5478
1116	CUUACAG G CCUUUCUA	1708	UAGAAAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUUGUAAG	5479
1126	UUUUCUA G UAAACAGU	1709	ACUGUUUA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUAGAAAAG	5480
1133	AGUAAACA G UAUUGAA	1710	UUCACAUU GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGUUUUACU	5481
1152	UUUACCCC G UUGCUCGG	1711	CCGAGCAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG GGGGUAAA	5482
1160	GUUGUUGG G CAACGGCC	1712	GGCCUUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CGAGCAAC	5483
1166	CGGCAACG G CCUGGUUC	1713	AGCCAGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CGUUGCCG	5484
1171	ACGGCUG G UCUAUGCC	1714	GGCAUAGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CAGGCCGU	5485
1182	UAUGCCAA G UGUUUGCU	1715	AGCAACA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUGGCAUA	5486

Table 42

1207	CCCCACUG G UUGGGGCU	1716	AGCCCCAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGUGGGG	5487
1213	UGGUUGGG G CUUGGCCA	1717	UGGCCAAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCAACCA	5488
1218	GGGGCUUG G CCAUAGGC	1718	GCCUAUGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAAGCCCC	5489
1225	GGCCAUAG G CCAUCAGC	1719	GCUGAUGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUAUGGCC	5490
1232	GGCCAUCA G CGCAUGCG	1720	CGCAUGCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGAUGGCC	5491
1240	GGCAUGC G UGGAACCU	1721	AGGUCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GCAUGCGC	5492
1287	AACUCCUA G CCGCUUGU	1722	ACAAGCGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UAGGAGUU	5493
1306	UGCUCGCA G CAGGUCUG	1723	CAGACCUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGCGAGCA	5494
1310	CGCAGCAG G UCUGGGGC	1724	GCCCCAGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGCUUGC	5495
1317	GGUCUGGG G CAAAACUC	1725	GAGUUUUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCAGACC	5496
1347	AUUCUUGC G UGCUCUCC	1726	GGAGAGCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GACAGAAU	5497
1379	UUUCCAU G CUGCUAGG	1727	CCUAGCAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAUGGAAA	5498
1387	GCUGCUAG G CUGUGCUG	1728	CAGCACAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUAGCAGC	5499
1418	CGCGGAC G UCCUUGU	1729	ACAAAGGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GUCCCGCG	5500
1431	UUGUJUAC G UCCCGUCG	1730	CGACGGGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GUAAACAA	5501
1436	UACGUCCC G UGGCGCU	1731	AGCGCCGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGGACGUA	5502
1440	UCCCGUCG G CGCUGAAU	1732	AUUCAGCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGACGGGA	5503
1471	CUCCCGGG G CCGCUUGG	1733	CCAAGCGG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCGGGAG	5504
1481	CGCUUGGG G CUCUACCG	1734	CGGUAGAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCAAGCG	5505
1517	UACCGACC G UCCACGGG	1735	CCCGUGGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGUCGGUA	5506
1526	UCCACGGG G CGCACCUC	1736	GAGGUCCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCGUGGA	5507
1553	GACUCCCC G UCUGUGCC	1737	GGCACAGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGGAGAGUC	5508
1579	GCCGGACC G UGUGCACU	1738	AGUGCACA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGUCCGGC	5509
1605	CUCUGCAC G UGCAUUGG	1739	CCAUGCGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GUGCAGAG	5510
1622	AGACCACC G UGAACGCC	1740	GGCGUUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GUUGGUCU	5511
1649	UGCCCAAG G UCUUGCAU	1741	AUGCAAGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGAAAGUC	5512
1679	GACUUUCA G CAUUGUCA	1742	UGACAUUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUUUGGCA	5513
1703	ACCUUGAG G CAUAUUUC	1743	GAAGUAUG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUCAAGGU	5514
1732	UUUAAUGA G UUGGAGGA	1744	UCCUCCCA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCAUUAAA	5515
1741	UGGGAGGA G UUGGGGGA	1745	UCCCCCAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCUCCCA	5516
1754	GGGAGGAG G UUAGGUUA	1746	UAACCUAA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUCCUCCC	5517
1759	GAGGUUAG G UUAAGGUU	1747	ACCUUUA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUAAACCUC	5518
1766	GGUUAAG G UCUUUGUA	1748	UACAAGA GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUUUAACC	5519
1782	ACUAGGAG G CUGUAGGC	1749	GCCUACAG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUCCUAGU	5520

Table 42

1789	GGCUGUAG G CAUAAAUU	1750	AAUUAUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUACAGCC	5521
1799	AUAAAUUG G UGUUUA	1751	UGAACACA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CAUUAU	5522
1811	GUUCACCA G CACCAUGC	1752	GCAUGGUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGGUGAAC	5523
1870	CUGUICAA G CCUCCAG	1753	CUUGGAGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUGAACAG	5524
1878	GCCUCCAA G CUGUGCCU	1754	AGGCACAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUGGAGGC	5525
1890	UGCCUUGG G UGGCUUUG	1755	CAAGGCCA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCAAGGCA	5526
1893	CUUGGGUG G CUUUGGGG	1756	CCCCAAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CACCCAAG	5527
1901	GCUTUGGG G CAUGGACA	1757	UGUCCAUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCCAAAGC	5528
1917	AUUGACCC G UAUAAAGA	1758	UCUTUAUA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG GGUCAAU	5529
1933	AAUTUGGA G CUUCUGUG	1759	CACAGAAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UCCAAAUU	5530
1944	UCUGUGGA G UUAUCUC	1760	GAGAGUAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UCCACAGA	5531
2023	AUCGGGGG G CCUUGAG	1761	CUCUAAGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCCCGAU	5532
2031	GCCUUGA G UCUCCGGA	1762	UCCGGAGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UCUAAGGC	5533
2062	ACCAUACG G CACUCAGG	1763	CCUGAGUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CGUAUGGU	5534
2070	GCACUCAG G CAAGCUAU	1764	AUAGCUUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUGAGUGC	5535
2074	UCAGGCAA G CUAUUCUG	1765	CAGAUAAG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUGCCUGA	5536
2090	GUGUUGGG G UGAGUUGA	1766	UCAACUCA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCCAACAC	5537
2094	UGGGUGGA G UUGAUGAA	1767	UUAUCUAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UCACCCCA	5538
2107	UGAAUCUA G CCACCUUG	1768	CCAGGUGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UAGAUAUA	5539
2116	CCACCUUG G UGGGAAGU	1769	ACUUCCCA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCAGGUGG	5540
2123	GGUGGGAA G UAAUUTUG	1770	CCAAUAUA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUCCACAC	5541
2140	AAGAUAUA G CAUCCAGG	1771	CCUGGAUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGGAUCUU	5542
2155	GGGAUAUA G UAGUCAGC	1772	GCUGACUA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UAAUUCUU	5543
2158	AAUAUGUA G UCAGCUAU	1773	AUAGCUUA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UACUAAUU	5544
2162	AGUAGUA G CUAUGUCA	1774	UGACUAUG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGACUAUU	5545
2173	AUGUCAC G UUAUAUUG	1775	CAUAUAUA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG GUUGACAU	5546
2183	UAAUAUGG G CUUAAAAA	1776	UUUUUAGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCUAUAUA	5547
2208	CUAUUGUG G UUUACAUU	1777	AUGUGAAA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CACAUAUG	5548
2235	ACUUUUGG G CGAGAAAC	1778	GUUUCUGG GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCAAAAGU	5549
2260	AAUAUUGG G UGUUUUUU	1779	AAAAGACA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CAAUAUUU	5550
2272	CUUUUGGA G UUGGUAUU	1780	AAUCCACA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UCCAAAAG	5551
2360	ACGAAGAG G CAGGUCCC	1781	GGGACCUU GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUUUUCGU	5552
2364	AGAGGCAG G UCCCCUAG	1782	CUAGGGGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUGCCUUA	5553
2403	AGACGAAG G UCUCUAUC	1783	GAUUGAGA GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUUUCGUU	5554

Table 42

2417	AUCGCCG G UGCAGAA	1784	UUCUGCGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG GCGGCGAU	5555
2454	CAAGUUA G UAUUCU	1785	AAGGAUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UAACAUUG	5556
2474	CACAUAG G UGGAAAC	1786	GUUCCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CUUAUGUG	5557
2491	UUUACGG G CUUAUUC	1787	GAUAAAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CCGUAAA	5558
2507	CUUCUAC G UACCUUG	1788	GCAAGGUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CGUAGAAG	5559
2530	CCUAAUG G CAACUCC	1789	GGAGUUUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CAUUUAGG	5560
2587	AGAUUA G CAUUUGU	1790	ACAAUUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UUAUAUCU	5561
2599	UUUGJGG G CCCCUIAC	1791	GUAAAGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CCCACAAA	5562
2609	CCCUUAC G UAAUUGAA	1792	UUCAUUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UGUAAAGG	5563
2650	CCUGCUAG G UUUUAUCC	1793	GGAUAAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CUAGCAGG	5564
2701	AUCAAAC G UAUAUCC	1794	GGAUUAUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG GGUUUGAU	5565
2713	UAUCCAGA G UAUGUAGU	1795	ACUAUAUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UCUGGAUA	5566
2720	AGUAUGUA G UUAUAU	1796	AUGAUUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UACAUACU	5567
2768	UUUGGAAG G CGGGAU	1797	GAUCCCG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CUUCCAAA	5568
2791	AAAGAGA G UCCACACG	1798	CGUGUGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UCUCUUUU	5569
2799	GUCCACAC G UAGCGCU	1799	AGGCGUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG GUGUGGAC	5570
2802	CACACGUA G CGCTUCAU	1800	AUGGCGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UACGUGUG	5571
2818	UUUGCGG G UCACCAUA	1801	UAUGUGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CCGCAAAA	5572
2848	GAUCUAC G CAUGGGAG	1802	CUCCAUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UGUAGAUC	5573
2857	CAUGGGAG G UUGGUCUU	1803	AAGACCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CUCCCAUG	5574
2861	GGAGGUUG G UCUCUCAA	1804	UUGGAUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CAACCUCC	5575
2881	UCGAAAG G AUGGGGA	1805	UCCCAUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CUUUUGA	5576
2936	GAUCAUA G UUGGACCC	1806	GGUCCAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UGAUGAUC	5577
2955	CAUUCAAA G CCAACUA	1807	UGAGUUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UUUGAAUG	5578
2964	GACAAUG G CCGGACG	1808	UGGAUUA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UGAGUUGG	5579
3005	CCAACAG G UGGAGUG	1809	GGUCCGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CAGUUGUC	5580
3021	AGUUGGA G UGGAGGUA	1810	CACUCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CUUGUUGG	5581
3027	AGUUGGA G UGGAGGUA	1811	UGCUCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UCCACCU	5582
3033	GAGUUGGA G CAUUCGG	1812	CCCGAUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UCCACUC	5583
3041	GCAUUCGG G CCAGGGU	1813	AACCUUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CCGAAUUG	5584
3047	GGGCCAGG G UUCACCC	1814	GGGUGAA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CCUGGCC	5585
3077	CUGUJGG G UGGAGCC	1815	GGGUCCA GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CCAACAG	5586
3082	GGGUGGA G CCCCACG	1816	CGUGAGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG UCCACCC	5587
3097	CGUCAGG G CUUACUA	1817	UGAGUAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCG CCUGAGCG	5588

Table 42

3117	CUGUGCCA G CAGCUCCU	1818	AGGAGCUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGGCACAG	5589
3120	UGCCAGCA G CUCCUCCU	1819	AGGAGGAG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGCUGGCA	5590
3146	ACCAUUG G CAGUCAGG	1820	CCUGACUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CGAUUGGU	5591
3149	AUUCGGCA G UCAGGAG	1821	CUUCCUGA GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGCCGAUU	5592
3158	UCAGGAAG G CAGCCUAC	1822	GUAGGCUG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CUUCCUGA	5593
3161	GGAAAGCA G CCUACUCC	1823	GGAGUAGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGCCUUC	5594
3204	AUCCUCAG G CCAUGCAG	1824	CUGCAUGG GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CUGAGGAU	5595
31	CUUUCUAA G AUCCGAGA	2196	UCUGGGAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UUGAAGAG	5596
38	AGAUCCCA G AGUCAGGG	2197	CCUGACU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGGGAUCU	5597
44	CAGAGUCA G GGCCUUGU	2198	ACAGGGCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGACUCUG	5598
45	AGAGUCAG G GCCCUGUA	2199	UACAGGGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CUGACUCU	5599
64	UUCUUGCU G GUGGCUCC	2200	GGAGCCAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGCAGGAA	5600
67	CUGCUGGU G GCUCGAGU	2201	ACUGGAGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACCAGCAG	5601
79	CCAGUUCA G GAACAGUG	2202	CACUGUUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGAACUGG	5602
80	CAGUUCAG G AACAGUGA	2203	UCACUGUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CUGAACUG	5603
99	CCUGCUCA G AAUACUGU	2204	ACAGUAUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGAGCAGG	5604
135	UUAUCGAA G ACUGGGGA	2205	UCCCCAGU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UUCGAUAA	5605
139	CGAAGACU G GGGACCCU	2206	AGGGUCCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AGUCUUCG	5606
140	GAAGACUG G GGACCCUG	2207	CAGGGUCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CAGUCUUC	5607
141	AAGACUGG G GACCCUGU	2208	ACAGGGUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CCAGUCUU	5608
142	AGACUGGG G ACCCUGUA	2209	UACAGGGU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CCCAGUCU	5609
159	CGAACAU G GAGAACAU	2210	AUGUUCUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUGUUCGG	5610
160	CGAACAU G GAGAACAU	2211	GAUGUUCU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CAUGUUCG	5611
162	AACAUUG G AACUCCUA	2212	GCGAUGUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UCCAUGUU	5612
175	UCGCAUCA G GATUCCUA	2213	UAGGAGUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGAUCCGA	5613
176	CGCAUCAG G ACUCCUAG	2214	CUAGGAGU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CUGAUGCG	5614
184	GACUCCUA G GACCCUUG	2215	CAGGGGUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UAGGAGUC	5615
185	ACUCCUAG G ACCCCUUG	2216	GCAGGGUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CUAGGAGU	5616
204	GUGUUA G GCGGGGUU	2217	AACCCCGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGUACAC	5617
207	UUACAGG G GGUUUUUU	2218	AAAAACCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG GCTUGUAA	5618
208	UACAGGG G GGUUUUUC	2219	GAAAAACC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CGCCUGUA	5619
209	ACAGGCG G GUUUUUUC	2220	AGAAAAAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CGCCUGU	5620
246	AUACCACA G AGUCUAGA	2221	UCUAGACU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UGUGGUUU	5621
253	AGAGUCUA G ACUCGUGG	2222	CCACGAGU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UAGACUCU	5622

Table 42

260	AGACUCGU G GUGACUU	2223	AAGUCCAC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG ACAGUCU	5623
263	CUCGUGGU G GACUUCUC	2224	GAGAAGUC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG ACCACGAG	5624
264	UCGUGGUG G ACUUCUCU	2225	AGAGAAGU GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CACCACGA	5625
283	AUUUUCUA G GGGAAACA	2226	UGUUCUCC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG UAGAAAUA	5626
284	UUUUCUAG G GGAACAC	2227	GUGUUCUCC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CUAGAAAA	5627
285	UUUCUAGG G GGAACACC	2228	GGUGUUCUCC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CCUAGAAA	5628
286	UUCUAGGG G GAACACCC	2229	GGGUGUUC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CCCUAGAA	5629
287	UCUAGGGG G AACACCCG	2230	CGGUGUUC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CCCUAGA	5630
304	UGUGUCUU G GCCAAAAU	2231	AUUUUGGC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG AAGACACA	5631
367	UUUGUCCU G GUUAUCGC	2232	GCGAUAC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG AGGACAAA	5632
377	UUUUGCU G GAUGUGUC	2233	GACACAUC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG AGCGAUAA	5633
378	UAUUGCUG G AUGUGUCU	2234	AGACACAU GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CAGCGAUA	5634
389	GUGUCUCG G GCGUUUUA	2235	UAAAACGC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG GCAGACAC	5635
441	UUCUUGUU G GUTUCUCU	2236	AGAAGAAC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG AACAGAAA	5636
450	GUUCUUCU G GACUAUCA	2237	UGAUAGUC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG AGAAGAAC	5637
451	UUCUUCUG G ACUAUCA	2238	UUGAUAGU GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CAGAAGAA	5638
460	ACUAUCA G GUUAUGUG	2239	CAACAUAU GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG UUGAUAGU	5639
490	UAUUUCCA G GAUCAUCA	2240	UGAUGAUC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG UGGAUUUA	5640
491	AAUUCACG G AUCAUCA	2241	UUGAUGAU GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CUGGAUUU	5641
511	CCAGCACG G GACCAUGC	2242	GCAUGGUC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG GGUGCUGG	5642
512	CAGCACCG G ACCAUGCA	2243	UGCAUGGU GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CGGUGCUG	5643
544	CUGCUCAA G GAACUCUCU	2244	AGAGGUUC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG UUGAGCAG	5644
545	UGCUCUAG G AACUCUCA	2245	UAGAGGUU GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CUUGAGCA	5645
585	AAACCUAC G GACGGAAA	2246	UUUCCGUC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG GUAGGUUU	5646
586	AACCUACG G ACGGAAAC	2247	GUUUCUGU GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CGUAGGUU	5647
589	CUACGGAC G GAAACUGC	2248	GCAGUUUC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG GUCCGUAG	5648
590	UACGGACG G AAACUGCA	2249	UGCAGUUU GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CGUCCGUA	5649
623	AUCAUCUU G GCUUUCUG	2250	CGAAAGCC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG AAGAUAU	5650
624	UCAUCUUG G GCUUUCGC	2251	GCGAAAGC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CAAGAUA	5651
644	AUACCUAU G GAGUGGGG	2252	CCCACUCC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG AUAGGUUU	5652
645	UACCUAUG G GAGUGGGC	2253	GCCCACUC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CAUAGGUA	5653
646	ACCUAUGG G AGUGGGGC	2254	GGCCCAUC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CCAUAGGU	5654
650	AUGGGAGU G GGCUCUAG	2255	CUGAGGCC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG ACUCCCAU	5655
651	UGGGAGUG G GCCUCAGU	2256	ACUGAGGC GGAGGAAACUCC CU UCAAGGACAUCGUCUCCGGG CACUCCCA	5656

Table 42

671	UUUCUCUU G GCUCAGUU	2257	AACUGAGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAGAGAAA	5657
701	UGUUCAGU G GUUCGUAG	2258	CUACGAAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG ACUGAACA	5658
709	GGUUCGUA G GGUUUUCC	2259	GGAAAGCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UACGAAAC	5659
710	GUUCGUAG G GCUUUUCC	2260	GGGAAAGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CUACGAAC	5660
728	CACUGUCU G GCUUUUCAG	2261	CUGAAAGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AGACAGUG	5661
743	AGUUAUUAU G GAUGAUGU	2262	ACAUCAUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUUAUAU	5662
744	GUUAUAUG G AUGAUGUG	2263	CACAUAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAUAUAAC	5663
752	GAUGAUGU G GUUUUUGG	2264	CCCAAAAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG ACAUCAUC	5664
758	GUGGUUUU G GGGGCCAA	2265	UUUGCCCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAAACCA	5665
759	UGGUUUUG G GGGCCAAG	2266	CUUGGCCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAAAACCA	5666
760	GGUUUUUG G GGCCAAGU	2267	ACUUGGCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAAAACCC	5667
761	GUUUUUGG G GCCAAGUC	2268	GACUUGGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CCCAAAAC	5668
824	UUGUCUUU G GGUUAACA	2269	UGUAUAAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAAGACAA	5669
825	UGUCUUUG G GUUAUACU	2270	AUGUAUAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAAAGACA	5670
856	AACAAAAA G AUGGGGAU	2271	AUCCCCAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UUUUUUUU	5671
859	AAAAAGAU G GGGUAUUU	2272	AUAUCCCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUCUUUUU	5672
860	AAAAGAUG G GGAUAUUC	2273	GAUAUUCG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAUCUUUU	5673
861	AAAAGAUG G GAUAUUCU	2274	GGAAUAUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CCAUCUUU	5674
862	AAGAUGGG G AUAUUCUCC	2275	GGGAUAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CCCAUCUU	5675
881	AACUUCAU G GGAUAUGU	2276	ACAUAUCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AUGAAGUU	5676
882	ACUUCAU G GAUAUGUA	2277	UACAUUUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAUGAAGU	5677
883	CUUCAUGG G AUAUGUA	2278	UUACAUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CCAUGAAG	5678
894	AUGUAUUU G GGAGUUGG	2279	CCAAUUCG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAUUAUACU	5679
895	UGUAUUUG G AGUUGGGG	2280	CCCAUUCG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAAUUAUAC	5680
896	GUUAUUUG G AGUUGGGG	2281	CCCAUUCG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CCAUUAUAC	5681
901	UGGAGUUU G GGGCACA	2282	AUGUGCCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AACUCCCA	5682
902	GGGAGUUU G GCACAUU	2283	AAUGUGCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CAACUCCC	5683
903	GGAGUUUG G GCACAUU	2284	CAUUGUCG GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CCAACUCC	5684
917	UUGCCACA G GAACAUU	2285	AUAUGUUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UGUGGCAA	5685
918	UGCCACAG G AACAUU	2286	AAUAUGUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CUGUGGCA	5686
952	GUGUUUUA G GAAAUUUC	2287	GAAGUUUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UAAAACAC	5687
953	UGUUUUAG G AAACUUC	2288	GGAGUUUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG CUAAAACA	5688
970	UGUAACA G GCCUAUUG	2289	CAUAAGGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG UGUUUACA	5689
982	UAUUGAUU G GAAAGUAU	2290	AUAUUUUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCCGG AAUCAUAU	5690

Table 42

983	AUUGAUUG G AAAGUAUG	2291	CAUACUUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAAUCAAU	5691
1004	CGAAUUGU G GGUCUUUU	2292	AAAAGACC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG ACAAUUCG	5692
1005	GAAUUGUG G GUCUUUUU	2293	CHAAAGAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CACAAUUC	5693
1013	GGUCUUUU G GGGUUUGC	2294	GCAAAACC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AAAAGACC	5694
1014	GUCUUUUG G GGUUUGCC	2295	GGCAAAACC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAAAAGAC	5695
1015	UCUUUUGG G GUUUGCCG	2296	CGGCAAAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCAAAAGA	5696
1041	CGCAUUGU G GAUAUUCU	2297	AGAAUAUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG ACAUUGCG	5697
1042	GCAUUGUG G AUAUUCUG	2298	CAGAAUAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CACAUTGC	5698
1088	GCAAAACA G GCUUUUAC	2299	GUAAAAGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGUUUUUGC	5699
1115	ACUUAACA G GCCUUUCU	2300	AGAAAGGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUGUAAGU	5700
1159	CGUUGCUC G GCAACGGC	2301	GCCGUUGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GAGCAACG	5701
1165	UCGGCAAC G GCCUGGUC	2302	GACCAGGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUUGCCGA	5702
1170	AACGGCCU G GUCUAUUG	2303	GCAUAGAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGGCCGUU	5703
1206	CCCCCACU G GUUGGGGC	2304	GCCCCAAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGUGGGGG	5704
1210	CACUGGUU G GGGCUTGG	2305	CCAAGCCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AACCAGUG	5705
1211	ACUGGUUG G GGCUUGGC	2306	GCCAAGCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAACCAGU	5706
1212	CUGGUUGG G GCUUGGCC	2307	GGCCAAGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCAACCAG	5707
1217	UGGGGCUU G GCCAUAGG	2308	CCUAUGGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AAGCCCCA	5708
1224	UGGCCAUA G GCCAUCAG	2309	CUGAUGGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UAUGGCCA	5709
1242	GCAUGCGU G GAACCUUU	2310	AAAGGUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG ACGCAUGC	5710
1243	CAUCCGUG G AACCUUUG	2311	CAAAGGUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CACGCAUG	5711
1277	AUACCGCG G AACUCUUA	2312	AGGAGUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GCGGUUUG	5712
1278	AUACCGCG G AACUCUUA	2313	UAGGAGUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CGCGGUUU	5713
1309	UCGCAGCA G GUCUGGGG	2314	CCCCAGAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGCUUGCGA	5714
1314	GCAGGUUU G GGGCAAAA	2315	UUUUGCCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGACCUUGC	5715
1315	CAGGUUCU G GGCAAAAC	2316	GUUUUGCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAGACCUUG	5716
1316	AGGUUCUG G GCAAAACU	2317	AGUUUUGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCAGACCU	5717
1329	AACUCAUC G GACUGAGC	2318	GUCAGUCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GAUGAGUU	5718
1330	ACUCAUCG G GACUGACA	2319	UGUCAGUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CGAUGAGU	5719
1331	CUCAUCGG G ACUGACAA	2320	UUGUCAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCGAUGAG	5720
1378	AUUUCCAU G GCUUGCUAG	2321	CUAGCAGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUGGAAAU	5721
1386	GGCUGCUA G GCUUGUCU	2322	AGCACAGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UAGCAGCC	5722
1402	UGCCAAAU G GAUCCUAC	2323	GUAGGAUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AGUUGGCA	5723
1403	GCCAACTUG G AUCCUACG	2324	CGUAGGAU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAGUUGGC	5724

Table 42

1413	UCCUACGC G GGACGUCC	2325	GGACGUCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GCGUAGGA	5725
1414	CCUACGGC G GACGUCCU	2326	AGGACGUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CGCGUAGG	5726
1415	CUACGGCG G ACGUCCUU	2327	AAGGACGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCGCGUAG	5727
1439	GUCCCGUC G GCGCUGAA	2328	UUCAGCGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GACGGGAC	5728
1454	AUCCCGCG G GACGACCC	2329	GGGUCGUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GCGGGAUU	5729
1455	AUCCCGCG G ACGACCCC	2330	GGGUCGUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GCGGGGAU	5730
1468	CCCUCCCG G GGGCGGCU	2331	AGCGGCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGGAGGGG	5731
1469	CCUCCCGG G GGCGGCUU	2332	AAGCGCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CGGGAGGG	5732
1470	CUCCCGCG G GCCGCUUG	2333	CAAGCGGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCGGGAGG	5733
1478	GGCGGCUU G GGGGCUUA	2334	UAGAGCCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AAGCGGCC	5734
1479	GCGGCUUG G GGCUCUAC	2335	GUAGAGCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAAGCGGC	5735
1480	CGGCUUGG G GCUUACCC	2336	GUAGAGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCAAGCGG	5736
1523	CGUCCAC G GGGGCGAC	2337	GUGCGCCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GUGGACGG	5737
1524	CGUCCACG G GGCACACC	2338	GUGCGCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CGUGGACG	5738
1525	GUCCACGG G GCGCACCU	2339	AGGUGCGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CCGUGGAC	5739
1544	CUUACGC G GACUCCCC	2340	GGGGAGUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GCGUAAAG	5740
1545	UUUACGGC G ACUCCCG	2341	CGGGGAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GCGUAAA	5741
1574	CAUCUGCC G GACCGUGU	2342	ACACGGUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGCAGAU	5742
1575	AUCUGCCG G ACCGUGUG	2343	CACACGGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG GGCAGAU	5743
1612	GUCGCAU G GAGACCAC	2344	GUGGUCUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AUGCGACG	5744
1613	GUCGAUG G AGACCACC	2345	GGUGUCU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAUGCGAC	5745
1615	CGCAUGA G ACCACCGU	2346	ACGGUGGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UCCAUGCG	5746
1635	CGCCACA G GAACCCUGC	2347	GCAGGUUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UGUGGGCG	5747
1636	GCCACAG G AACCGGCC	2348	GGCAGGUU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CUGUGGGC	5748
1648	CUGCCCAA G GUCUUGCA	2349	UGCAAGAC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUGGCGAG	5749
1660	UUGCAUAA G AGGACUCU	2350	AGAGUCU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUAUGCAA	5750
1662	GCAUAGA G GACUCUUG	2351	CAAGAGUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UCUUAUGC	5751
1663	CAUAGAG G ACUCUUGG	2352	CCAAGAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CUCUUAUG	5752
1670	GGACUCUU G GACUUUCA	2353	UGAAAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG AAGAGUCC	5753
1671	GACUCUUG G ACUUUCAG	2354	CUGAAAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CAAGAGUC	5754
1702	GACCUUGA G GCAUACUU	2355	AAGUAGC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UCAAGGUC	5755
1715	ACUCCAAA G ACUGUGUG	2356	CACACAGU GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG UUGAAGU	5756
1734	UAUGAGU G GGAGGAGU	2357	ACUCCUCC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG ACUCAUA	5757
1735	AUAGAGU G GAGGAGUU	2358	AACUCCUC GGAGGAAACUCC CU UCAAGGACAUCGUCCGGG CACUCAUU	5758

Table 42

1736	AUGAGUGG G AGGAGUUG	2359	CAACUCCU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CCACUCAU	5759
1738	GAGUGGGA G GAGUTUGG	2360	CCCAACUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UCCACACU	5760
1739	AGUGGGAG G AGUUGGGG	2361	CCCCAACU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CUCCACU	5761
1744	GAGGAGUU G GGGGAGGA	2362	UCCUCCCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AACUCCUC	5762
1745	AGGAGUUU G GGGAGGAG	2363	CUCCUCCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CAACUCCU	5763
1746	GGAGUUGG G GGAGGAGG	2364	CCUCCUCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CCAACUCC	5764
1747	GAGUUGGG G GAGGAGGU	2365	ACCUCUCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CCCAACUC	5765
1748	AGUUGGGG G AGGAGGUU	2366	AACUCCU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CCCAACU	5766
1750	UUGGGGGA G GAGGUUAG	2367	CURACCUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UCCGCCAA	5767
1751	UGGGGGAG G AGGUUAGG	2368	CCUACCUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CUCCGCCA	5768
1753	GGGGGGA G GUUAGGUU	2369	AACCUAAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UCCUCCCC	5769
1758	GGAGGUUA G GUUAAAGG	2370	CCUUUAAAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UAACUCCU	5770
1765	AGUUAAA G GUCUUUGU	2371	ACAAAGAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UUUAACCU	5771
1778	UUGUACUA G GAGGCUGU	2372	ACAGCCUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UAGUACAA	5772
1779	UGUACUAG G AGGCTUGA	2373	UACAGCCU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CUAGUACA	5773
1781	UACUAGGA G GCUGUAGG	2374	CCUACAGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UCCUAGUA	5774
1788	AGGCUGUA G GCAUAAAU	2375	AUUUAUGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UACAGCCU	5775
1798	CAUAAAUU G GUGUGUUC	2376	GAACACAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAUUUAUG	5776
1888	UGUGCCUU G GGUGGCUU	2377	AAGCCACC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAGGCACA	5777
1889	GUGCCUUG G GUGGCUUU	2378	AAAGCCAC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CAAAGGCAC	5778
1892	CCUUGGGU G GCUUUGGG	2379	CCCAAGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACCCAAGG	5779
1898	GUGGCUUU G GGGCAUGG	2380	CCAUGCCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAAGCCAC	5780
1899	UGGCUUUG G GCAUGGAC	2381	UCCAUGCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CAAAGCCA	5781
1900	GGCUUUGG G GCAUUGA	2382	GUCCAUGC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CCAAGGCC	5782
1905	UGGGGCAU G GACAUUGA	2383	UCAUUGUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AUGCCCCA	5783
1906	GGGGCAUG G ACAUUGAC	2384	GUCAUUGU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CAUGCCCC	5784
1924	CGUAUAAA G AAUUUGGA	2385	UCCAAAUU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UUUAUACG	5785
1930	AAGAAUUU G GAGCUUCU	2386	AGAAGCUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG AAUUAUCU	5786
1931	AGAAUUUG G AGCUUCUG	2387	CAGAAGCU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CAAAUUCU	5787
1941	GCUCUCUG G GAGUUAACU	2388	AGUAACUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG ACAGAAAGC	5788
1942	CUUCUCUG G AGUUAACUC	2389	GAGUACUC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CACAGAAG	5789
1987	CUAUUGGA G AUCUCCUC	2390	GAGGAGAU GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG UCGAAUAG	5790
2018	UCUGUAUC G GGGGGCCU	2391	AGGCCCCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG GAUACAGA	5791
2019	CUGUAUCG G GGGGCCUU	2392	AAGGCCCC GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG CGAUACAG	5792

Table 42

2020	UGUAUCGG G GGGCCUUA	2393	UAAGGCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCGAUACA	5793
2021	GUUAUCGG G GGCUUAG	2394	CUAAGGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCGAUAC	5794
2022	UAUCGGGG G GCCUUAGA	2395	UCUAAGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCCGAUA	5795
2029	GGGCCUUA G AGUCUCCG	2396	CGGAGCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UAAGGCC	5796
2037	GAGUCUC G GAACAUUG	2397	CAAUUGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GGAGACUC	5797
2038	AGUCUCCG G AACAUUGU	2398	ACAAUUGU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCGAGACU	5798
2061	CACCAUAC G GCACUCAG	2399	CUGAGUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GUAUGGUG	5799
2069	GGCACAUA G GCAAGCUA	2400	UAGTUUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGAGUGCC	5800
2087	UCUGUGU G GGUUGAGU	2401	ACUCACC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AACACAGA	5801
2088	CUGUGUUG G GGUUGAGU	2402	AACUCACC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAACACAG	5802
2089	UGUGUUGG G GUGAGUUG	2403	CAACUCAC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCAACACA	5803
2114	AGCCACCU G GUGGGAA	2404	UUCACC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AGUGGCU	5804
2115	GCCACCUG G GUGGGAAG	2405	CUUCCAC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAGGUGGC	5805
2118	ACCUGGU G GGAAGUAA	2406	UUACUCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACCCAGGU	5806
2119	CCUGGGUG G GAAGUAAU	2407	AUUAUUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CACCAGG	5807
2120	CUGGGUGG G AAGUAAU	2408	AAUUAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCACCCAG	5808
2130	AGUAAUU G GAAGAUCC	2409	GGAUUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAUUAU	5809
2131	GUAAUUUG G AAGAUCUA	2410	UGGAUUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAAAUUAC	5810
2134	AUUUGGAA G AUCCAGCA	2411	UGCUGAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUCCAAU	5811
2147	AGCAUCCA G GGAUUUAG	2412	CUAAUUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGGAUGCU	5812
2148	GCAUCCAG G GAUUUAGU	2413	ACUAAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGGAUGC	5813
2149	CAUCCAGG G AAUUUAGUA	2414	UACUAAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUUGGAUG	5814
2181	GUUAAUAU G GGCUAAA	2415	UUUAGCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUUAUUAAC	5815
2182	UUAUAUG G GCUAAA	2416	UUUAGGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAUAUUA	5816
2195	AAAAUCA G ACAACUAU	2417	AUAGUUGU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGAUUUUU	5817
2207	ACUAUUUG G GUUUCACA	2418	UGUAAAC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACAUAGU	5818
2233	UAUUUUU G GCGAGAA	2419	UUCUCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAAAGUA	5819
2234	UACUUUG G GCGAGAA	2420	UUUCCG GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAAAGUA	5820
2239	UGGGCGA G AAACUGUU	2421	AACAUUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCGCCAA	5821
2259	GAUAUUU G GUGUCUUU	2422	AAAGAC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAUAUUC	5822
2269	UGUCUUU G GAGUGUG	2423	CCACAC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAAAGACA	5823
2270	GUCUUUG G AGUGUGGA	2424	UCCACU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAAAGAC	5824
2276	UGGAGUGU G GAUUCGCA	2425	UGGAAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG ACACUCCA	5825
2277	GGAGUGUG G AUUCGCAC	2426	GUGGAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CACACUCC	5826

Table 42

2300	UGCAUUA G ACCACCAA	2427	UUGUGGU GGAGAAACUCC CU UCAAGGACAUCGUCGGG UAUAUGCA	5827
2334	ACACUCC G GAACUAC	2428	GUAGUUC GGAGAAACUCC CU UCAAGGACAUCGUCGGG GGAAGUGU	5828
2335	CACUCCG G AAACUACU	2429	AGUAGUU GGAGAAACUCC CU UCAAGGACAUCGUCGGG CGAAGUG	5829
2351	UGUUGUUA G ACGAAGAG	2430	CUCUCCU GGAGAAACUCC CU UCAAGGACAUCGUCGGG UAACACAA	5830
2357	UAGACGAA G AGGAGGU	2431	ACCUGCCU GGAGAAACUCC CU UCAAGGACAUCGUCGGG UUCGUCUA	5831
2359	GACGAAG G GAGGUCC	2432	GGACUUC GGAGAAACUCC CU UCAAGGACAUCGUCGGG UCUUGGUC	5832
2363	AAGAGGA G GUCCCCUA	2433	UAGGGAC GGAGAAACUCC CU UCAAGGACAUCGUCGGG UGCCUCUU	5833
2372	GUCCCCUA G AAGAAGAA	2434	UUCUUCU GGAGAAACUCC CU UCAAGGACAUCGUCGGG UAGGGAC	5834
2375	CCCUAGAA G AAGAACUC	2435	GAGUUCU GGAGAAACUCC CU UCAAGGACAUCGUCGGG UUCUAGGG	5835
2378	UAGAAGAA G AACUCCU	2436	AGGAGUU GGAGAAACUCC CU UCAAGGACAUCGUCGGG UUCUUCUA	5836
2396	GCCUCGA G ACGAAGGU	2437	ACCUCCU GGAGAAACUCC CU UCAAGGACAUCGUCGGG UGCAGGC	5837
2402	CAGACGAA G GUCUCAU	2438	AUTGAGC GGAGAAACUCC CU UCAAGGACAUCGUCGGG UUCGUCUG	5838
2423	GCGUCGA G AAGAUCUC	2439	GAGAUUU GGAGAAACUCC CU UCAAGGACAUCGUCGGG UGCAGCG	5839
2426	UCGAGAA G AUCUCAU	2440	AUTGAGU GGAGAAACUCC CU UCAAGGACAUCGUCGGG UUCUGCGA	5840
2438	UCAAUCU G GAAUCUC	2441	GAGAUUC GGAGAAACUCC CU UCAAGGACAUCGUCGGG GAGAUUGA	5841
2439	CAAUCCG G GAAUCUA	2442	UGAGAUU GGAGAAACUCC CU UCAAGGACAUCGUCGGG CGAGAUUG	5842
2440	AAUCCCG G AAUCUCA	2443	UUGAGAU GGAGAAACUCC CU UCAAGGACAUCGUCGGG CCGAGAUU	5843
2463	UAUCCUU G GACACUA	2444	UAUGUUC GGAGAAACUCC CU UCAAGGACAUCGUCGGG AAGGAUA	5844
2464	AUCCUUG G ACACUAU	2445	UAUUGUU GGAGAAACUCC CU UCAAGGACAUCGUCGGG CAAGGAU	5845
2473	ACACUAU G GUGGAAA	2446	UUUCCAC GGAGAAACUCC CU UCAAGGACAUCGUCGGG UUAUGUGU	5846
2476	CAUAAGU G GAAACUU	2447	AAGUUUC GGAGAAACUCC CU UCAAGGACAUCGUCGGG ACCUUAUG	5847
2477	AUAAGGU G GAAACUU	2448	AAAGUUU GGAGAAACUCC CU UCAAGGACAUCGUCGGG CACCUUAU	5848
2478	UAAGGUG G AAACUUU	2449	UAAUUUU GGAGAAACUCC CU UCAAGGACAUCGUCGGG CCACCUUA	5849
2488	AACUUAC G GGCUUUA	2450	UAAAGCC GGAGAAACUCC CU UCAAGGACAUCGUCGGG GUAAAGUU	5850
2489	ACUUUAC G GGCUUUA	2451	AUAAAGC GGAGAAACUCC CU UCAAGGACAUCGUCGGG CGUAAAGU	5851
2490	CUUUACG G GCUUUUU	2452	AAUAAAG GGAGAAACUCC CU UCAAGGACAUCGUCGGG CCGUAAAG	5852
2506	UCUUCUAC G GUACCUU	2453	CAAGUAC GGAGAAACUCC CU UCAAGGACAUCGUCGGG GUAGAAGA	5853
2529	UCCUAAU G GCAAACUC	2454	GAGUUUC GGAGAAACUCC CU UCAAGGACAUCGUCGGG AUUUAGGA	5854
2563	CAUUUGCA G GAGGACAU	2455	AUGUCCU GGAGAAACUCC CU UCAAGGACAUCGUCGGG UGCAAAUG	5855
2564	AUUUGCAG G AGGACAUU	2456	AUGUCCU GGAGAAACUCC CU UCAAGGACAUCGUCGGG CUGCAAAU	5856
2566	UUGCAGG G ACUAUUGU	2457	ACAAUUC GGAGAAACUCC CU UCAAGGACAUCGUCGGG UCCUGCAA	5857
2567	UGCAGGAG G ACUAUUGU	2458	AACAUGU GGAGAAACUCC CU UCAAGGACAUCGUCGGG CUCCUGCA	5858
2580	UGUUGAU G AUGUAAGC	2459	GCTUACU GGAGAAACUCC CU UCAAGGACAUCGUCGGG UAUCACAA	5859
2596	CAUUUUGU G GGGCCCU	2460	AGGGCCC GGAGAAACUCC CU UCAAGGACAUCGUCGGG ACAAAUUG	5860

Table 42

2597	AAUUGUG G GGCCCUU	2461	AAGGGCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CACAAAUU	5861
2598	AUUUGUG G GCCCUUA	2462	UAAGGGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCACAAU	5862
2622	UGAAACA G GAGACUA	2463	UAAGUCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGUUUUA	5863
2623	GAAACAG G AGACUAA	2464	UUAAGUCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUGUUUC	5864
2625	AAACAGG A ACUUAUU	2465	AUUUAAGU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCUGUUU	5865
2649	GCCUGCUA G GUUUUAU	2466	GAUAAAC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UAGCAGGC	5866
2684	UGCCCUUA G AUAAGGG	2467	CCCUUAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UAAGGCA	5867
2690	UAGUAAA G GAUCAAA	2468	UUUGAUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUUUAUCU	5868
2691	AGAUAAAG G GAUCAAC	2469	GUUGAUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CUUUUAUCU	5869
2692	GAUAAAG G AUCAAAC	2470	GUUUGAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCUUUAUC	5870
2711	AUUAUCCA G AGUAUGUA	2471	UACUAACU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGGAUAAU	5871
2737	UACUCCA G ACGGACA	2472	UGUGCGU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UGGAAGUA	5872
2763	CACUCUUU G GAAGGCG	2473	CCGCCUUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAAGAGUG	5873
2764	ACUCUUUG G AAGGCGG	2474	CCGCCUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAAAGAGU	5874
2767	CUUUGGA G GCGGGAU	2475	AUCCCGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUCCAAAG	5875
2770	UGAAGGC G GGAUCUU	2476	AAGAUCU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GCCUCCA	5876
2771	GGAAGCG G GGAUCUA	2477	UAAGAUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGCCUCC	5877
2772	GAAGCGG G GAUCUUA	2478	AUAAGAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCGCCUUC	5878
2773	AAGGCGG G AUCUUAU	2479	UAUAAAU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCGCCUU	5879
2787	AUAUAAA G AGAGUCCA	2480	UGGACUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUUUAUU	5880
2789	AUAAAGA G AGUCCACA	2481	UGUGACU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCUUUUU	5881
2816	CAUUUUG G GGUCCACA	2482	UGUGACC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG GCACAAUG	5882
2817	AUUUGCG G GUCACCAU	2483	AUGGUGAC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CGCAAAU	5883
2832	AUAUUCU G GGAACAAG	2484	CUUGUUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AAGAAUU	5884
2833	UAUUCUG G GAACAAGA	2485	UCUUGUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAAGAAU	5885
2834	AUCUUGG G AACAAGAU	2486	AUCUUGU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCAAGAAU	5886
2840	GGGAACA G AUCUACAG	2487	CUGUAUU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUGUUCU	5887
2852	ACAGCAU G GGAGUUG	2488	CAACUCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUGCUGUA	5888
2853	ACAGCAU G GAGGUUG	2489	CCAACUC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CAUGCUGU	5889
2854	CAGCAUG G AGGUUGU	2490	ACCAACU GGAGAAACUCC CU UCAAGGACAUCGUCCGGG CCAUGCUG	5890
2856	GCAUGGA G GUUGUCU	2491	AGACCAAC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UCCCAUGC	5891
2860	GGGAGGU G GUCUCCA	2492	UGGAGAC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AACCUCC	5892
2880	CUCGAAA G GCAUGGG	2493	CCCAUGC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG UUUUCGAG	5893
2885	AAAGCAU G GGGACAA	2494	UUUGUCC GGAGAAACUCC CU UCAAGGACAUCGUCCGGG AUGCCUUU	5894

Table 42

2886	AAGGCAUG G GGACAAAU	2495	AUUUGUCC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CAUGCCUU	5895
2887	AGGCAUGG G GACAAAU	2496	GAUUUGC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CCAUGCCU	5896
2888	GGCAUGGG G ACAAUUCU	2497	GAUUUGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CCAUGCC	5897
2915	AUCCCCU G GAUUCUU	2498	AAGAAUCC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AGGGGAU	5898
2916	AUCCCCU G GAUUCUUC	2499	GAAGAAUC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CAGGGGAU	5899
2917	UCCCCUUG G AUUCUUC	2500	GAAGAAU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CCAGGGGA	5900
2939	CAUCAGUU G GACCCUGC	2501	GCAGGUU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AACUGAU	5901
2940	AUCAGUUG G ACCCUGCA	2502	UGCAGGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CAACUGAU	5902
2973	UAAAUCCA G AUUGGGAC	2503	GUCCAAU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UGGAUUA	5903
2977	UCCAGAUU G GGACCUCA	2504	UGAGGUU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AAUCUGGA	5904
2978	CCAGAUUG G GACCUCAA	2505	UUGAGGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CAAUCUGG	5905
2979	CAGAUUG G ACCUCAAC	2506	GUUGAGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CCAUCUG	5906
2996	CCGCACAA G GACAAACU	2507	CAGUUGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UTUGCGG	5907
2997	CGCACAA G GACAAACU	2508	CCAGUUGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CUUGUGCG	5908
3004	GGACAACU G GCGGACG	2509	CGUCCGC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AGUUGUCC	5909
3008	AACUGGCG G GACGCCAA	2510	UUGGCUU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG GGCCAGU	5910
3009	ACUGGCG G ACGCCAAC	2511	GUUGGCU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CGGCCAGU	5911
3020	GCCAAACA G GUGGGAGU	2512	ACUCCAC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UUGUUGGC	5912
3023	AACAAGU G GAGUGGG	2513	CCACUCC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG ACCUUGU	5913
3024	ACAAGGU G GAGUGGGA	2514	UCCACU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CACCUUGU	5914
3025	CAAGGU G AGUGGGAG	2515	CUCCACU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CCACCUUG	5915
3029	GUGGGAGU G GGAGCAU	2516	AAUGCU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG ACUCCAC	5916
3030	UGGGAGU G GAGCAUUC	2517	GAUUGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CACUCCCA	5917
3031	GGGAGU G AGCAUUCG	2518	CGAAUGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CACUCC	5918
3039	GAGCAUUC G GGCCAGGG	2519	CCUUGGC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG GAAUGCU	5919
3040	AGCAUUC G GCCAGGGU	2520	ACCUUGC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CGAAUGCU	5920
3045	UCGGGCA G GUUCACC	2521	GGUAAAC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG UGGCCCGA	5921
3046	CGGGCCAG G GUUCACC	2522	GGUAAAC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CUGGCCCG	5922
3063	UCCCAU G GGGACU	2523	CAGUCC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AUGGGGAG	5923
3064	UCCCAU G GGGACU	2524	ACAGUCC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CAUGGGGA	5924
3065	CCCAUG G GGACU	2525	AACAGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CCAUGGG	5925
3066	CCCAUG G GACU	2526	CAACAGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CCAUGGG	5926
3067	CCAUGGG G ACUGU	2527	CCAAGU GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG CCCAUGG	5927
3074	GGACUUGU G GGGUGG	2528	CUCCACC GGAGGAAACUCC CU UCAAGGACAUUGUCCGGG AACAGUCC	5928

Table 42

3075	GACUGUUG G GGUGGAGC	2529	GCUCCACC GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CAACAGUC	5929
3076	ACUGUUGG G GUGGAGCC	2530	GGCUCCAC GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CCAACAGU	5930
3079	GUUGGGGU G GAGCCUC	2531	GAGGGCUC GGAGAAACUCC CU UCAAGGACAUCGUCGCGG ACCCCAAC	5931
3080	UUGGGGUG G AGCCUCA	2532	UGAGGGCU GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CACCCCAA	5932
3095	CAGGCUA G GGCUACU	2533	AGUAGGCC GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGAGCGUG	5933
3096	ACGUCAG G GCCUACUC	2534	GAGUAGGC GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUGAGCGU	5934
3145	CACCAUC G GCAGUCAG	2535	CUGACUGC GGAGAAACUCC CU UCAAGGACAUCGUCGCGG GAUUGGUG	5935
3153	GGCAGUA G GAAGGCAG	2536	CUGCCUUC GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGACUGCC	5936
3154	GCAGUCAG G AAGGCAGC	2537	GCUGCCUU GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUGACUGC	5937
3157	GUAGGAA G GCAGCCUA	2538	UAGGCUUC GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUCUUGAC	5938
3187	ACCUCUAA G GGACACUC	2539	GAGUUCU GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UUAGAGGU	5939
3188	CCUCUAG G GACACUCA	2540	UGAGUUC GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUUAGAGG	5940
3189	CUCUAAG G ACACUCAU	2541	AUGAGUGU GGAGAAACUCC CU UCAAGGACAUCGUCGCGG CUCUAGAG	5941
3203	CAUCCUCA G GCCAUGCA	2542	UGCAUGGC GGAGAAACUCC CU UCAAGGACAUCGUCGCGG UGAGGAUG	5942

Input Sequence = AF100308. Cut Site = YG/M or UG/U.

Stem Length = 8. Core Sequence = GGAGGAAACUCC CU UCAAGGACAUCGUCGCGG
 AF100308 (Hepatitis B virus strain 2-18, 3215 bp)

Table 43

Table 43: Human HBV Ribozyme and Target Sequence

Pos	Substrate	Seq ID	RP#	Ribozyme Alias	Ribozyme	Rz Seq ID
313	CCAAAAU U CGCAGUC	5943	18157	HBV-313 Rz-7 RNA	GACUGCG CUGAUGAGGCCCGUAGGCCGAA AUUUUGG B	6293
327	CCCAAAU C UCCAGUC	5944	18158	HBV-327 Rz-7 RNA	GACUGGA CUGAUGAGGCCCGUAGGCCGAA AUUUUGG B	6294
334	CUCCAGU C ACUCACC	5945	18159	HBV-334 Rz-7 RNA	GGUGAGU CUGAUGAGGCCCGUAGGCCGAA ACUGGAG B	6295
408	UCUUCU C UGCAUCC	5946	18160	HBV-408 Rz-7 RNA	GGAUGCA CUGAUGAGGCCCGUAGGCCGAA AGGAAGA B	6296
557	UCUAUGU U UCCCUCA	5947	18161	HBV-557 Rz-7 RNA	UGAGGGA CUGAUGAGGCCCGUAGGCCGAA ACAUAGA B	6297
1255	UUUGUGU C UCCUCUG	5948	18162	HBV-1255 Rz-7 RNA	CAGAGGA CUGAUGAGGCCCGUAGGCCGAA ACACAAA B	6298
1538	CCUCUCU U UACGGCG	5949	18163	HBV-1538 Rz-7 RNA	CCGCGUA CUGAUGAGGCCCGUAGGCCGAA AGAGAGG B	6299
1756	AGGAGGU U AGGUUAA	5950	18164	HBV-1756 Rz-7 RNA	UUAACCU CUGAUGAGGCCCGUAGGCCGAA ACCUCCU B	6300
1861	AUGUCCU A CUGUACA	5951	18165	HBV-1861 Rz-7 RNA	UGAACAG CUGAUGAGGCCCGUAGGCCGAA AGGACAU B	6301
2504	UUCUUCU A CGGUACC	5952	18166	HBV-2504 Rz-7 RNA	GGUACCG CUGAUGAGGCCCGUAGGCCGAA AGAAGAA B	6302
10	CUCCACC A CUUUCUA	5953	18197	HBV-10 CHz-7 RNA	UGGAAAG CUGAUGAGGCCCGUAGGCCGAA GGUGGAG B	6303
335	UCCAGUC A CUCACCA	5954	18198	HBV-335 CHz-7 RNA	UGGUGAG CUGAUGAGGCCCGUAGGCCGAA GACUGGA B	6304
1258	GUGUCUC C UCUGCCG	5955	18199	HBV-1258 CHz-7 RNA	CGGCAGA CUGAUGAGGCCCGUAGGCCGAA GAGACAC B	6305
2307	GACCACC A AAUGCCC	5956	18200	HBV-2307 CHz-7 RNA	GGGCAUU CUGAUGAGGCCCGUAGGCCGAA GGUGGUC B	6306
347	UCACCAACCU G UUGUC	5957	18216	HBV-347 GCI.Rz-5/10 RNA	GACAA UGAUGGCAUGCACUAUGCGCG AGGUUGGUGA B	6307
350	CCAACCCUUU G UCCUC	5958	18217	HBV-350 GCI.Rz-5/10 RNA	GAGGA UGAUGGCAUGCACUAUGCGCG AACAGGUUGG B	6308
1508	UCCGCCUAUU G UACCG	5959	18218	HBV-1508 GCI.Rz-5/10 RNA	CGGUA UGAUGGCAUGCACUAUGCGCG AAUAGGCGGA B	6309
234	AAUCCU C ACAUA	5960	18334	HBV-234 Rz-6 allyl stab1	u ₅ -a ₅ -u ₅ -gu cUGAuGagggccguuagggccGaa Aggaau B	6310
252	GAGUCU A GACUCG	5961	18335	HBV-252 Rz-6 allyl stab1	c ₅ g ₅ a ₅ g ₅ uc cUGAuGagggccguuagggccGaa Agacuc B	6311
268	UGGACU U CUCUCA	5962	18337	HBV-268 Rz-6 allyl stab1	u ₅ g ₅ a ₅ g ₅ ag cUGAuGagggccguuagggccGaa Agucca B	6312
280	AAUUUU C UAGGGG	5963	18345	HBV-280 Rz-6 allyl stab1	c ₅ c ₅ c ₅ c ₅ ua cUGAuGagggccguuagggccGaa Aaaaau B	6313
313	CAAAAU U CGCAGU	5964	18346	HBV-313 Rz-6 allyl stab1	a ₅ c ₅ u ₅ g ₅ cg cUGAuGagggccguuagggccGaa Auuuug B	6314
395	GGCGUU U UAUCAU	5965	18350	HBV-395 Rz-6 allyl stab1	a ₅ u ₅ g ₅ a ₅ ua cUGAuGagggccguuagggccGaa Aacgcc B	6315
402	UAUCAU C UCCUC	5966	18351	HBV-402 Rz-6 allyl stab1	g ₅ -a ₅ g ₅ g ₅ aa cUGAuGagggccguuagggccGaa Augaua B	6316
607	UGUAUU C CCAUCC	5967	18355	HBV-607 Rz-6 allyl stab1	g ₅ g ₅ a ₅ u ₅ gg cUGAuGagggccguuagggccGaa Auauca B	6317
697	UUUGUU C AGUGGU	5968	18362	HBV-697 Rz-6 allyl stab1	a ₅ c ₅ c ₅ a ₅ cu cUGAuGagggccguuagggccGaa Aacaaa B	6318
1539	UCUCUU U ACGCGG	5969	18366	HBV-1539 Rz-6 allyl stab1	c ₅ s ₅ g ₅ c ₅ sgu cUGAuGagggccguuagggccGaa Aagaga B	6319
1599	UCACCU C UGCACG	5970	18367	HBV-1599 Rz-6 allyl stab1	c ₅ g ₅ u ₅ g ₅ ca cUGAuGagggccguuagggccGaa Agguga B	6320
1607	GCACGU C GCAUGG	5971	18368	HBV-1607 Rz-6 allyl stab1	c ₅ -c ₅ -a ₅ -u ₅ -gc cUGAuGagggccguuagggccGaa Acgugc B	6321

Table 43

1833	UCACCU C UGCCUA	5972	18371	HBV-1833 Rz-6 allyl stab1	u ₅ a ₅ g ₅ s ₅ ca cUGAuGagggccguuagggccGaa Agguga B	6322
2383	AGAACU C CCUGGC	5973	18374	HBV-2383 Rz-6 allyl stab1	g ₅ c ₅ g ₅ a ₅ s ₅ gg cUGAuGagggccguuagggccGaa Aguuu B	6323
2429	GAAGAU C UCAUCC	5974	18376	HBV-2429 Rz-6 allyl stab1	g ₅ a ₅ s ₅ u ₅ u ₅ sga cUGAuGagggccguuagggccGaa Auuuuc B	6324
2831	UAUUCU U GGGAAAC	5975	18379	HBV-2831 Rz-6 allyl stab1	g ₅ u ₅ s ₅ c ₅ s ₅ cc cUGAuGagggccguuagggccGaa Agaaau B	6325
430	UGCCUC A UCUUCU	5976	18391	HBV-430 CHz-6 allyl stab1	a ₅ g ₅ a ₅ s ₅ ga cUGAuGagggccguuagggccGaa laggca B	6326
676	UGGCUC A GUUUAC	5977	18396	HBV-676 CHz-6 allyl stab1	g ₅ u ₅ a ₅ s ₅ ac cUGAuGagggccguuagggccGaa lagcca B	6327
683	GUUUAC U AGUGCC	5978	18397	HBV-683 CHz-6 allyl stab1	g ₅ g ₅ c ₅ a ₅ s ₅ cu cUGAuGagggccguuagggccGaa luaaac B	6328
1150	UUIUAC C CGUUGC	5979	18402	HBV-1150 CHz-6 allyl stab1	g ₅ c ₅ a ₅ s ₅ c ₅ cg cUGAuGagggccguuagggccGaa lguaaa B	6329
1200	GCAACC C CCACUG	5980	18403	HBV-1200 CHz-6 allyl stab1	c ₅ a ₅ s ₅ u ₅ sgg cUGAuGagggccguuagggccGaa lguuuc B	6330
1201	CAACCC C CACUGG	5981	18404	HBV-1201 CHz-6 allyl stab1	c ₅ c ₅ a ₅ g ₅ ug cUGAuGagggccguuagggccGaa lggguug B	6331
1444	CGGCGC U GAAUCC	5982	18405	HBV-1444 CHz-6 allyl stab1	g ₅ g ₅ a ₅ u ₅ uc cUGAuGagggccguuagggccGaa lgcucc B	6332
1451	GAAUCC C GCGGAC	5983	18406	HBV-1451 CHz-6 allyl stab1	g ₅ u ₅ c ₅ s ₅ gc cUGAuGagggccguuagggccGaa lgaauuc B	6333
1533	CGCACC U CUCUUU	5984	18407	HBV-1533 CHz-6 allyl stab1	a ₅ a ₅ s ₅ g ₅ ag cUGAuGagggccguuagggccGaa lgugcg B	6334
1600	CACCUC U GCACGU	5985	18410	HBV-1600 CHz-6 allyl stab1	a ₅ c ₅ g ₅ u ₅ sgc cUGAuGagggccguuagggccGaa laggug B	6335
1698	CCGACC U UGAGGC	5986	18411	HBV-1698 CHz-6 allyl stab1	g ₅ c ₅ s ₅ u ₅ ca cUGAuGagggccguuagggccGaa lguccg B	6336
1784	GGAGGC U GUAGGC	5987	18412	HBV-1784 CHz-6 allyl stab1	g ₅ c ₅ s ₅ u ₅ ac cUGAuGagggccguuagggccGaa lcuucc B	6337
1829	UUUUUC A CCUCUG	5988	18414	HBV-1829 CHz-6 allyl stab1	c ₅ a ₅ g ₅ a ₅ gg cUGAuGagggccguuagggccGaa laaaaa B	6338
1876	GCCUCC A AGCUGU	5989	18420	HBV-1876 CHz-6 allyl stab1	a ₅ c ₅ a ₅ g ₅ s ₅ cu cUGAuGagggccguuagggccGaa lgaggc B	6339
1880	CCAAGC U GUGCCU	5990	18422	HBV-1880 CHz-6 allyl stab1	a ₅ g ₅ g ₅ c ₅ ac cUGAuGagggccguuagggccGaa lcuugg B	6340
218	UUUUUC U GUUGACA	5991	18333	HBV-218 Rz-7 allyl stab1	u ₅ g ₅ u ₅ c ₅ aac cUGAuGagggccguuagggccGaa Agaaaa B	6341
257	CUAGACU C GUGGUGG	5992	18336	HBV-257 Rz-7 allyl stab1	c ₅ c ₅ a ₅ c ₅ ac cUGAuGagggccguuagggccGaa Agucuag B	6342
268	GUGGACU C UCUCUAA	5993	18338	HBV-268 Rz-7 allyl stab1	u ₅ g ₅ a ₅ gag cUGAuGagggccguuagggccGaa Aguccac B	6343
269	UGGACUU C UCUCAAU	5994	18339	HBV-269 Rz-7 allyl stab1	a ₅ u ₅ u ₅ g ₅ aga cUGAuGagggccguuagggccGaa Aaguoca B	6344
271	GACUUCU C UCAAUUU	5995	18340	HBV-271 Rz-7 allyl stab1	a ₅ a ₅ a ₅ u ₅ uga cUGAuGagggccguuagggccGaa Agaaguc B	6345
273	CUUCUCU C AAUUUUC	5996	18341	HBV-273 Rz-7 allyl stab1	g ₅ a ₅ a ₅ auu cUGAuGagggccguuagggccGaa Aungag B	6346
277	UCUCAAU U UUCUAGG	5997	18342	HBV-277 Rz-7 allyl stab1	c ₅ c ₅ u ₅ a ₅ gaa cUGAuGagggccguuagggccGaa Aungaga B	6347
278	CUCAAUU U UCUAGGG	5998	18343	HBV-278 Rz-7 allyl stab1	c ₅ c ₅ s ₅ u ₅ aga cUGAuGagggccguuagggccGaa Aauugag B	6348
279	UCAAUUU U CUAGGGG	5999	18344	HBV-279 Rz-7 allyl stab1	c ₅ c ₅ c ₅ u ₅ ag cUGAuGagggccguuagggccGaa Aaauga B	6349
314	CAAAUUU C GCAGUCC	6000	18347	HBV-314 Rz-7 allyl stab1	g ₅ g ₅ a ₅ c ₅ ugc cUGAuGagggccguuagggccGaa Aauuug B	6350
385	GAUGUGU C UGCGGCG	6001	18348	HBV-385 Rz-7 allyl stab1	c ₅ g ₅ s ₅ c ₅ gca cUGAuGagggccguuagggccGaa Acatauc B	6351
394	GCGGCGU U UUAUCAU	6002	18349	HBV-394 Rz-7 allyl stab1	a ₅ u ₅ g ₅ a ₅ uaa cUGAuGagggccguuagggccGaa Acgccc B	6352

Table 43

402	UUAUCAU C UUCCUCU	6003	18352	HBV-402 Rz-7 allyl stab1	a ₃ g ₃ a ₃ g ₃ gaa cUGAUgagccguuagccGaa Augauaa B	6353
423	UGCUGCU A UGCCUCA	6004	18353	HBV-423 Rz-7 allyl stab1	u ₃ g ₃ a ₃ g ₃ gca cUGAUgagccguuagccGaa Agcagca B	6354
429	UAUGCCU C AUCUUCU	6005	18354	HBV-429 Rz-7 allyl stab1	a ₃ g ₃ a ₃ a ₃ gau cUGAUgagccguuagccGaa Aggcaua B	6355
679	GCUCAGU U UACUAGU	6006	18356	HBV-679 Rz-7 allyl stab1	a ₃ g ₃ u ₃ a ₃ gua cUGAUgagccguuagccGaa Acugagc B	6356
680	CUCAGUU U ACUAGUG	6007	18357	HBV-680 Rz-7 allyl stab1	c ₃ a ₃ c ₃ u ₃ agu cUGAUgagccguuagccGaa Aacugag B	6357
681	UCAGUUU A CUAGUGC	6008	18358	HBV-681 Rz-7 allyl stab1	g ₃ c ₃ a ₃ c ₃ uag cUGAUgagccguuagccGaa Aaacuga B	6358
684	GUUUAU A GUGCCAU	6009	18359	HBV-684 Rz-7 allyl stab1	a ₃ u ₃ g ₃ g ₃ cac cUGAUgagccguuagccGaa Aguaaac B	6359
692	GUGCCAU U UGUUCAG	6010	18360	HBV-692 Rz-7 allyl stab1	c ₃ u ₃ g ₃ a ₃ aca cUGAUgagccguuagccGaa Auggcac B	6360
693	UGCCAUU U GUUCAGU	6011	18361	HBV-693 Rz-7 allyl stab1	a ₃ c ₃ u ₃ g ₃ aac cUGAUgagccguuagccGaa Auuggca B	6361
1534	CGCACCU C UCUUUC	6012	18363	HBV-1534 Rz-7 allyl stab1	g ₃ u ₃ a ₃ a ₃ aga cUGAUgagccguuagccGaa Aggugcg B	6362
1536	CACCUUC C UUUACGC	6013	18364	HBV-1536 Rz-7 allyl stab1	g ₃ c ₃ g ₃ u ₃ aaa cUGAUgagccguuagccGaa Agaggug B	6363
1538	CCUCUCU U UACGCGG	6014	18365	HBV-1538 Rz-7 allyl stab1	c ₃ c ₃ g ₃ c ₃ gua cUGAUgagccguuagccGaa Agagagg B	6364
1787	AGGCUGU A GGCAUAA	6015	18369	HBV-1787 Rz-7 allyl stab1	u ₃ u ₃ a ₃ u ₃ g ₃ cc cUGAUgagccguuagccGaa Acagccu B	6365
1793	UAGGCAU A AAUUGGU	6016	18370	HBV-1793 Rz-7 allyl stab1	a ₃ c ₃ c ₃ a ₃ auu cUGAUgagccguuagccGaa Augccua B	6366
1874	CAAGCCU C CAAGCUG	6017	18372	HBV-1874 Rz-7 allyl stab1	c ₃ a ₃ g ₃ c ₃ uug cUGAUgagccguuagccGaa Aggcuug B	6367
1887	UGUGCCU U GGGUGGC	6018	18373	HBV-1887 Rz-7 allyl stab1	g ₃ c ₃ c ₃ a ₃ ccc cUGAUgagccguuagccGaa Aggcaca B	6368
2383	AAGAAU C CCUCGCC	6019	18375	HBV-2383 Rz-7 allyl stab1	g ₃ g ₃ c ₃ g ₃ agg cUGAUgagccguuagccGaa Aguuuuu B	6369
2828	ACCAUUAU C UUGGGGA	6020	18377	HBV-2828 Rz-7 allyl stab1	u ₃ c ₃ c ₃ g ₃ aag cUGAUgagccguuagccGaa Auauuggu B	6370
2829	CCAUAUU C UUGGGAA	6021	18378	HBV-2829 Rz-7 allyl stab1	u ₃ u ₃ c ₃ c ₃ caa cUGAUgagccguuagccGaa Aauauagg B	6371
2831	AUAUUCU U GGGAACA	6022	18380	HBV-2831 Rz-7 allyl stab1	u ₃ g ₃ u ₃ u ₃ ccc cUGAUgagccguuagccGaa Agauuuu B	6372
256	UCUAGAC U CGUGGUG	6023	18381	HBV-256 CHZ-7 allyl stab1	c ₃ a ₃ c ₃ c ₃ acg cUGAUgagccguuagccGaa luuaga B	6373
267	GGUGGAC U UCUCUCA	6024	18382	HBV-267 CHZ-7 allyl stab1	u ₃ g ₃ a ₃ g ₃ aga cUGAUgagccguuagccGaa luuccacc B	6374
270	GGACUUC U CUCAAUU	6025	18383	HBV-270 CHZ-7 allyl stab1	a ₃ a ₃ u ₃ u ₃ gag cUGAUgagccguuagccGaa laagucc B	6375
272	ACUUCUC U CAUUUUU	6026	18384	HBV-272 CHZ-7 allyl stab1	a ₃ a ₃ a ₃ u ₃ g cUGAUgagccguuagccGaa lagaagu B	6376
274	UUCUCUC A AUUUUCU	6027	18385	HBV-274 CHZ-7 allyl stab1	a ₃ g ₃ a ₃ a ₃ auu cUGAUgagccguuagccGaa lagagaa B	6377
386	AUGUGUC U GCGGCGU	6028	18386	HBV-386 CHZ-7 allyl stab1	a ₃ c ₃ g ₃ c ₃ g ₃ cc cUGAUgagccguuagccGaa lacacau B	6378
419	AUCCUGC U GCUAUGC	6029	18387	HBV-419 CHZ-7 allyl stab1	g ₃ c ₃ a ₃ u ₃ agc cUGAUgagccguuagccGaa lcaggau B	6379
422	CUGCUGC U AUGCCUC	6030	18388	HBV-422 CHZ-7 allyl stab1	g ₃ a ₃ g ₃ g ₃ cau cUGAUgagccguuagccGaa lcaggag B	6380
427	GCUAUGC C UCAUCUU	6031	18389	HBV-427 CHZ-7 allyl stab1	a ₃ a ₃ g ₃ a ₃ uga cUGAUgagccguuagccGaa lcauagc B	6381
428	CUAUGCC U CAUCUUC	6032	18390	HBV-428 CHZ-7 allyl stab1	g ₃ a ₃ a ₃ g ₃ aug cUGAUgagccguuagccGaa lgcauag B	6382
430	AUGCCUC A UCUUCUU	6033	18392	HBV-430 CHZ-7 allyl stab1	a ₃ a ₃ g ₃ a ₃ aga cUGAUgagccguuagccGaa laggcau B	6383

Table 43

608	UGUAUUC C CAUCCCA	6034	18393	HBV-608 CHZ-7 allyl stab1	u ₅ g ₅ g ₅ aug cUGAuGagggccguaggccGaa laauaca B	6384
609	GUUUUCC C AUCCCAU	6035	18394	HBV-609 CHZ-7 allyl stab1	a ₅ u ₅ g ₅ g ₅ gau cUGAuGagggccguaggccGaa lgaauac B	6385
669	GUUUCUC U UGGCUCU	6036	18395	HBV-669 CHZ-7 allyl stab1	u ₅ g ₅ a ₅ g ₅ cca cUGAuGagggccguaggccGaa lgaauac B	6386
689	CUAGUGC C AUUUGUU	6037	18398	HBV-689 CHZ-7 allyl stab1	a ₅ a ₅ c ₅ a ₅ aa cUGAuGagggccguaggccGaa lcauag B	6387
690	UAGUGCC A UUUGUUC	6038	18399	HBV-690 CHZ-7 allyl stab1	g ₅ a ₅ a ₅ c ₅ aaa cUGAuGagggccguaggccGaa lgcacua B	6388
718	GUUUUCC C CCACUGU	6039	18400	HBV-718 CHZ-7 allyl stab1	a ₅ c ₅ a ₅ g ₅ ugg cUGAuGagggccguaggccGaa lgaauag B	6389
1149	CCUUUAC C CCGUUGC	6040	18401	HBV-1149 CHZ-7 allyl stab1	g ₅ c ₅ a ₅ a ₅ cgg cUGAuGagggccguaggccGaa luuaagg B	6390
1535	GCACCUC U CUUUACG	6041	18408	HBV-1535 CHZ-7 allyl stab1	c ₅ g ₅ u ₅ a ₅ ag cUGAuGagggccguaggccGaa laggugc B	6391
1537	ACCUCUC U UUACGCG	6042	18409	HBV-1537 CHZ-7 allyl stab1	c ₅ g ₅ c ₅ g ₅ uaa cUGAuGagggccguaggccGaa lagaggu B	6392
1791	UGUAGGC A UAAAUUG	6043	18413	HBV-1791 CHZ-7 allyl stab1	c ₅ a ₅ a ₅ u ₅ uaa cUGAuGagggccguaggccGaa lccuaca B	6393
1831	UUUUCAC C UCUGCCU	6044	18415	HBV-1831 CHZ-7 allyl stab1	a ₅ g ₅ g ₅ c ₅ aga cUGAuGagggccguaggccGaa lugaana B	6394
1832	UUUCACC U CUGCCUA	6045	18416	HBV-1832 CHZ-7 allyl stab1	u ₅ a ₅ g ₅ g ₅ cag cUGAuGagggccguaggccGaa lgugaaa B	6395
1872	UUCAAGC C UCCAAGC	6046	18417	HBV-1872 CHZ-7 allyl stab1	g ₅ c ₅ u ₅ g ₅ gga cUGAuGagggccguaggccGaa lcuugaa B	6396
1873	UCAAGCC U CCAAGCU	6047	18418	HBV-1873 CHZ-7 allyl stab1	a ₅ g ₅ c ₅ u ₅ ugg cUGAuGagggccguaggccGaa lgcuaa B	6397
1875	AAGCCUC C AAGCUGU	6048	18419	HBV-1875 CHZ-7 allyl stab1	a ₅ c ₅ a ₅ g ₅ cua cUGAuGagggccguaggccGaa laggcuu B	6398
1876	AGCCUCC A AGCUGUG	6049	18421	HBV-1876 CHZ-7 allyl stab1	c ₅ a ₅ c ₅ a ₅ gcu cUGAuGagggccguaggccGaa laggcuu B	6399
1880	UCCAAGC U GUGCCUU	6050	18423	HBV-1880 CHZ-7 allyl stab1	a ₅ a ₅ g ₅ g ₅ cac cUGAuGagggccguaggccGaa lcuugga B	6400
2382	GAAGAAC U CCCUCGC	6051	18424	HBV-2382 CHZ-7 allyl stab1	g ₅ c ₅ g ₅ a ₅ g ₅ g cUGAuGagggccguaggccGaa luucuuu B	6401
2384	AGAACUC C CUCGCCU	6052	18425	HBV-2384 CHZ-7 allyl stab1	a ₅ g ₅ g ₅ c ₅ gag cUGAuGagggccguaggccGaa laggcuu B	6402
2385	GAACUCC C UCGCCUC	6053	18426	HBV-2385 CHZ-7 allyl stab1	g ₅ a ₅ g ₅ g ₅ cga cUGAuGagggccguaggccGaa laggcuu B	6403
2422	GCGUCCG A GAAGAUC	6054	18427	HBV-2422 CHZ-7 allyl stab1	g ₅ g ₅ u ₅ c ₅ uuc cUGAuGagggccguaggccGaa lcgacgc B	6404
2830	CAUAUUC U UGGGAAC	6055	18428	HBV-2830 CHZ-7 allyl stab1	g ₅ u ₅ u ₅ c ₅ cca cUGAuGagggccguaggccGaa laauaug B	6405
234	AAUCCU C ACAUA	6056	19179	HBV-234 Rz-6 amino stab1	u ₅ a ₅ u ₅ g ₅ gu cUGAuGagggccguaggccGaa Aggaau B	6406
252	GAGUCU A GACUCG	6057	19180	HBV-252 Rz-6 amino stab1	c ₅ g ₅ a ₅ g ₅ uc cUGAuGagggccguaggccGaa Agacuc B	6407
268	UGGACU U CUCUCA	6058	19182	HBV-268 Rz-6 amino stab1	u ₅ g ₅ a ₅ g ₅ ag cUGAuGagggccguaggccGaa Agucua B	6408
280	AAUUUU C UAGGGG	6059	19190	HBV-280 Rz-6 amino stab1	c ₅ c ₅ c ₅ ua cUGAuGagggccguaggccGaa Aaaauu B	6409
313	CAAAAU U CGCAGU	6060	19191	HBV-313 Rz-6 amino stab1	a ₅ c ₅ u ₅ g ₅ c cUGAuGagggccguaggccGaa Auuuug B	6410
395	GGCGUU U UAUCAU	6061	19195	HBV-395 Rz-6 amino stab1	a ₅ u ₅ g ₅ a ₅ ua cUGAuGagggccguaggccGaa Aagcc B	6411
402	UAUCAU C UUCCUC	6062	19196	HBV-402 Rz-6 amino stab1	g ₅ a ₅ g ₅ g ₅ aa cUGAuGagggccguaggccGaa Augaua B	6412
607	UGUAUU C CCAUCC	6063	19200	HBV-607 Rz-6 amino stab1	g ₅ g ₅ a ₅ u ₅ g ₅ g cUGAuGagggccguaggccGaa Auuaa B	6413
697	UUUGUU C AGUGGU	6064	19207	HBV-697 Rz-6 amino stab1	a ₅ c ₅ c ₅ a ₅ c cUGAuGagggccguaggccGaa Aacaaa B	6414

Table 43

1539	UCUCUU U ACGCGG	6065	19211	HBV-1539 Rz-6 amino stab1	c ₅ s ₅ s ₅ c ₅ gu cUGAUGaggccguuaggccGaa Aagaga B	6415
1599	UCACCU C UGCACG	6066	19212	HBV-1599 Rz-6 amino stab1	c ₅ s ₅ u ₅ s ₅ ca cUGAUGaggccguuaggccGaa Agguga B	6416
1607	GCACGU C GCAUGG	6067	19213	HBV-1607 Rz-6 amino stab1	c ₅ s ₅ a ₅ u ₅ gc cUGAUGaggccguuaggccGaa Acgugc B	6417
1833	UCACCU C UGCCUA	6068	19216	HBV-1833 Rz-6 amino stab1	u ₅ s ₅ s ₅ s ₅ ca cUGAUGaggccguuaggccGaa Agguga B	6418
2383	AGAACU C CCUCGC	6069	19219	HBV-2383 Rz-6 amino stab1	g ₅ s ₅ s ₅ a ₅ gg cUGAUGaggccguuaggccGaa Aguuc B	6419
2429	GAAGAU C UCAAUC	6070	19221	HBV-2429 Rz-6 amino stab1	g ₅ a ₅ u ₅ u ₅ ga cUGAUGaggccguuaggccGaa Aucuuc B	6420
2831	UAUUCU U GGAAC	6071	19224	HBV-2831 Rz-6 amino stab1	g ₅ u ₅ u ₅ c ₅ cc cUGAUGaggccguuaggccGaa Agaaua B	6421
430	UGCCUC A UCUUCU	6072	19236	HBV-430 CHZ-6 amino stab1	a ₅ s ₅ a ₅ s ₅ ga cUGAUGaggccguuaggccGaa laggca B	6422
676	UGGCUC A GUUUAC	6073	19241	HBV-676 CHZ-6 amino stab1	g ₅ u ₅ a ₅ s ₅ ac cUGAUGaggccguuaggccGaa lagcca B	6423
683	GUUUAC U AGUGCC	6074	19242	HBV-683 CHZ-6 amino stab1	g ₅ s ₅ c ₅ a ₅ cu cUGAUGaggccguuaggccGaa luaaac B	6424
1150	UUUACC C CGUUGC	6075	19247	HBV-1150 CHZ-6 amino stab1	g ₅ c ₅ a ₅ a ₅ cg cUGAUGaggccguuaggccGaa lguaaa B	6425
1200	GCAACC C CCACUG	6076	19248	HBV-1200 CHZ-6 amino stab1	c ₅ a ₅ g ₅ u ₅ gg cUGAUGaggccguuaggccGaa lguugc B	6426
1201	CAACCC C CACUGG	6077	19249	HBV-1201 CHZ-6 amino stab1	c ₅ c ₅ a ₅ g ₅ ug cUGAUGaggccguuaggccGaa lggug B	6427
1444	CGGCGC U GAAUCC	6078	19250	HBV-1444 CHZ-6 amino stab1	g ₅ s ₅ a ₅ u ₅ uc cUGAUGaggccguuaggccGaa lgcgcg B	6428
1451	GAAUCC C GCGGAC	6079	19251	HBV-1451 CHZ-6 amino stab1	g ₅ u ₅ c ₅ s ₅ gc cUGAUGaggccguuaggccGaa lgauc B	6429
1533	CGCACC U CUCUUU	6080	19252	HBV-1533 CHZ-6 amino stab1	a ₅ a ₅ a ₅ s ₅ ag cUGAUGaggccguuaggccGaa lgugcg B	6430
1600	CACCUC U GCACGU	6081	19255	HBV-1600 CHZ-6 amino stab1	a ₅ c ₅ g ₅ u ₅ gc cUGAUGaggccguuaggccGaa laggug B	6431
1698	CCGACC U UGAGGC	6082	19256	HBV-1698 CHZ-6 amino stab1	g ₅ c ₅ c ₅ u ₅ ca cUGAUGaggccguuaggccGaa lgucgg B	6432
1784	GGAGGC U GUAGGC	6083	19257	HBV-1784 CHZ-6 amino stab1	g ₅ c ₅ s ₅ u ₅ ac cUGAUGaggccguuaggccGaa lccucc B	6433
1829	UUUUUC A CCUCUG	6084	19259	HBV-1829 CHZ-6 amino stab1	c ₅ a ₅ s ₅ a ₅ gg cUGAUGaggccguuaggccGaa laaaaa B	6434
1876	GCCUCC A AGCUGU	6085	19265	HBV-1876 CHZ-6 amino stab1	a ₅ c ₅ a ₅ s ₅ cu cUGAUGaggccguuaggccGaa lgaggc B	6435
1880	CCAAGC U GUGCCU	6086	19267	HBV-1880 CHZ-6 amino stab1	a ₅ s ₅ s ₅ c ₅ ac cUGAUGaggccguuaggccGaa lcuugg B	6436
218	UUUUUCU U GUUGACA	6087	19178	HBV-218 Rz-7 amino stab1	u ₅ s ₅ u ₅ c ₅ aac cUGAUGaggccguuaggccGaa Agaaaa B	6437
257	CUAGACU C GUGGUGG	6088	19181	HBV-257 Rz-7 amino stab1	c ₅ c ₅ a ₅ c ₅ cac cUGAUGaggccguuaggccGaa Agucua B	6438
268	GUGGACU U CUCUCA	6089	19183	HBV-268 Rz-7 amino stab1	u ₅ u ₅ g ₅ a ₅ gag cUGAUGaggccguuaggccGaa Aguccac B	6439
269	UGGACUU C UCUCAU	6090	19184	HBV-269 Rz-7 amino stab1	a ₅ u ₅ u ₅ s ₅ aga cUGAUGaggccguuaggccGaa Aagucca B	6440
271	GACUUCU C UCAAUUU	6091	19185	HBV-271 Rz-7 amino stab1	a ₅ a ₅ a ₅ u ₅ ga cUGAUGaggccguuaggccGaa Agaaguc B	6441
273	CUUCUCU C AAUUUUC	6092	19186	HBV-273 Rz-7 amino stab1	g ₅ a ₅ a ₅ auu cUGAUGaggccguuaggccGaa Agagaag B	6442
277	UCUCAU U UUCUAGG	6093	19187	HBV-277 Rz-7 amino stab1	c ₅ c ₅ u ₅ a ₅ gaa cUGAUGaggccguuaggccGaa Auugaga B	6443
278	CUCAAUU U UCUAGGG	6094	19188	HBV-278 Rz-7 amino stab1	c ₅ s ₅ c ₅ u ₅ aga cUGAUGaggccguuaggccGaa Aauugag B	6444
279	UCAAUUU U CUAGGGG	6095	19189	HBV-279 Rz-7 amino stab1	c ₅ s ₅ c ₅ s ₅ uag cUGAUGaggccguuaggccGaa Aauuuga B	6445

Table 43

314	CAAAUU C GCAGUCC	6096	19192	HBV-314 Rz-7 amino stab1	g ₉ s ₉ a ₉ c ₉ ugc cUGAUGagccguuagccGaa Auuuug B	6446
385	GAUGUG C UGCGGCG	6097	19193	HBV-385 Rz-7 amino stab1	c ₉ s ₉ c ₉ s ₉ gca cUGAUGagccguuagccGaa Acacauc B	6447
394	GCGGCGU U UUAUCAU	6098	19194	HBV-394 Rz-7 amino stab1	a ₉ u ₉ s ₉ a ₉ uaa cUGAUGagccguuagccGaa Acgccgc B	6448
402	UUAUCAU C UUCCUCU	6099	19197	HBV-402 Rz-7 amino stab1	a ₉ s ₉ a ₉ s ₉ gaa cUGAUGagccguuagccGaa Augaua B	6449
423	UGCUGCU A UGCCUCA	6100	19198	HBV-423 Rz-7 amino stab1	u ₉ s ₉ a ₉ g ₉ gca cUGAUGagccguuagccGaa Agcagca B	6450
429	UAUGOCU C AUCUUCU	6101	19199	HBV-429 Rz-7 amino stab1	a ₉ s ₉ a ₉ s ₉ gaa cUGAUGagccguuagccGaa Aggcaua B	6451
679	GCUCAGU U UACUAGU	6102	19201	HBV-679 Rz-7 amino stab1	a ₉ s ₉ u ₉ s ₉ gua cUGAUGagccguuagccGaa Acugagc B	6452
680	CUCAGUU U ACUAGUG	6103	19202	HBV-680 Rz-7 amino stab1	c ₉ s ₉ c ₉ u ₉ agu cUGAUGagccguuagccGaa Aacuga B	6453
681	UCAGUUU A CUAGUGC	6104	19203	HBV-681 Rz-7 amino stab1	g ₉ s ₉ a ₉ s ₉ uag cUGAUGagccguuagccGaa Auaaac B	6454
684	GUUUACU A GUGCCAU	6105	19204	HBV-684 Rz-7 amino stab1	a ₉ u ₉ s ₉ s ₉ cac cUGAUGagccguuagccGaa Auggac B	6455
692	GUGCCAU U GUUCAGU	6106	19205	HBV-692 Rz-7 amino stab1	c ₉ u ₉ s ₉ a ₉ aca cUGAUGagccguuagccGaa Aaagc B	6456
693	UGCCACU C UCUUUAU	6107	19206	HBV-693 Rz-7 amino stab1	a ₉ s ₉ u ₉ s ₉ aac cUGAUGagccguuagccGaa Auugc B	6457
1534	CGCACCU C UCUUUAU	6108	19208	HBV-1534 Rz-7 amino stab1	g ₉ u ₉ a ₉ s ₉ aga cUGAUGagccguuagccGaa Aggucg B	6458
1536	CACCUCU C UUUACGC	6109	19209	HBV-1536 Rz-7 amino stab1	g ₉ s ₉ s ₉ u ₉ aaa cUGAUGagccguuagccGaa Agaggu B	6459
1538	CCUCUCU U UACGCGG	6110	19210	HBV-1538 Rz-7 amino stab1	c ₉ s ₉ g ₉ c ₉ gua cUGAUGagccguuagccGaa Agaggg B	6460
1787	AGGCUGU A GGCAUAA	6111	19214	HBV-1787 Rz-7 amino stab1	u ₉ u ₉ a ₉ u ₉ gcc cUGAUGagccguuagccGaa Acagccu B	6461
1793	UAGGCAU A AAUUGGU	6112	19215	HBV-1793 Rz-7 amino stab1	a ₉ s ₉ c ₉ a ₉ uuu cUGAUGagccguuagccGaa Augcua B	6462
1874	CAAGCCU C CAAGCUG	6113	19217	HBV-1874 Rz-7 amino stab1	c ₉ a ₉ g ₉ s ₉ uug cUGAUGagccguuagccGaa Aggcuug B	6463
1887	UGUGCCU U GGGUGGC	6114	19218	HBV-1887 Rz-7 amino stab1	g ₉ s ₉ c ₉ a ₉ ccc cUGAUGagccguuagccGaa Aggcaca B	6464
2383	AAGAACU C CCUCGCC	6115	19220	HBV-2383 Rz-7 amino stab1	g ₉ s ₉ c ₉ g ₉ agg cUGAUGagccguuagccGaa Aguuuu B	6465
2828	ACCAUUAU U CUUGGGA	6116	19222	HBV-2828 Rz-7 amino stab1	u ₉ s ₉ c ₉ s ₉ aag cUGAUGagccguuagccGaa Auauggu B	6466
2829	CCAUUAU C UUGGGAA	6117	19223	HBV-2829 Rz-7 amino stab1	u ₉ u ₉ c ₉ s ₉ caa cUGAUGagccguuagccGaa Auaugg B	6467
2831	AUAUUCU U GGGAAACA	6118	19225	HBV-2831 Rz-7 amino stab1	u ₉ g ₉ u ₉ u ₉ ccc cUGAUGagccguuagccGaa Agaauau B	6468
256	UCUAGAC U CGUGGUG	6119	19226	HBV-256 CHz-7 amino stab1	c ₉ a ₉ c ₉ s ₉ acg cUGAUGagccguuagccGaa luuaga B	6469
267	GGUGGAC U UCUCUCA	6120	19227	HBV-267 CHz-7 amino stab1	u ₉ g ₉ a ₉ g ₉ aga cUGAUGagccguuagccGaa luccacc B	6470
270	GGACUUC U CUCAAUU	6121	19228	HBV-270 CHz-7 amino stab1	a ₉ a ₉ u ₉ u ₉ gag cUGAUGagccguuagccGaa laagu B	6471
272	ACUUCUC U CAAUUIU	6122	19229	HBV-272 CHz-7 amino stab1	a ₉ a ₉ a ₉ u ₉ uug cUGAUGagccguuagccGaa lagaagu B	6472
274	UUCUCUC A AUUIUCU	6123	19230	HBV-274 CHz-7 amino stab1	a ₉ s ₉ a ₉ a ₉ aau cUGAUGagccguuagccGaa lagagaa B	6473
386	AUGUGUC U GCGGCGU	6124	19231	HBV-386 CHz-7 amino stab1	a ₉ c ₉ s ₉ c ₉ gcg cUGAUGagccguuagccGaa lacacau B	6474
419	AUCCUGC U GCUAUGC	6125	19232	HBV-419 CHz-7 amino stab1	g ₉ s ₉ a ₉ u ₉ agc cUGAUGagccguuagccGaa lcaggau B	6475
422	CUGCUGC U AUGCCUC	6126	19233	HBV-422 CHz-7 amino stab1	g ₉ a ₉ g ₉ s ₉ cau cUGAUGagccguuagccGaa lcagcag B	6476

Table 43

427	GCUAUGC C UCAUCUU	6127	19234	HBV-427 CHZ-7 amino stab1	a ₅ a ₅ g ₅ a ₅ uga cUGAUgagccgcuuagccGaa lcauagc B	6477
428	CUAUGCC U CAUCUUC	6128	19235	HBV-428 CHZ-7 amino stab1	g ₅ a ₅ a ₅ g ₅ aug cUGAUgagccgcuuagccGaa lgaucag B	6478
430	AUGCCUC A UCUCUU	6129	19237	HBV-430 CHZ-7 amino stab1	a ₅ a ₅ g ₅ a ₅ aga cUGAUgagccgcuuagccGaa laggcau B	6479
608	UGUAUUC C CAUCCCA	6130	19238	HBV-608 CHZ-7 amino stab1	u ₅ g ₅ g ₅ g ₅ aug cUGAUgagccgcuuagccGaa laauaca B	6480
609	GUUAUUC C AUCCCAU	6131	19239	HBV-609 CHZ-7 amino stab1	a ₅ u ₅ g ₅ g ₅ gau cUGAUgagccgcuuagccGaa lgaauac B	6481
669	GUUUCUC U UGGCUCA	6132	19240	HBV-669 CHZ-7 amino stab1	u ₅ g ₅ a ₅ g ₅ cca cUGAUgagccgcuuagccGaa lgaauac B	6482
689	CUAGUGC C AUUUGUU	6133	19243	HBV-689 CHZ-7 amino stab1	a ₅ a ₅ g ₅ a ₅ aa cUGAUgagccgcuuagccGaa lcaucag B	6483
690	UAGUGCC A UUUUGUUC	6134	19244	HBV-690 CHZ-7 amino stab1	g ₅ a ₅ a ₅ g ₅ aaa cUGAUgagccgcuuagccGaa lgcacua B	6484
718	GCUUUCC C CCACUGU	6135	19245	HBV-718 CHZ-7 amino stab1	a ₅ c ₅ a ₅ g ₅ ugg cUGAUgagccgcuuagccGaa lgaagc B	6485
1149	CCUUUAC C CCGUUGC	6136	19246	HBV-1149 CHZ-7 amino stab1	g ₅ c ₅ a ₅ a ₅ cgg cUGAUgagccgcuuagccGaa luuaagg B	6486
1535	GCACCUU C CUUUACG	6137	19253	HBV-1535 CHZ-7 amino stab1	c ₅ g ₅ u ₅ a ₅ agg cUGAUgagccgcuuagccGaa laggugc B	6487
1537	ACCUCUC U UUACGCG	6138	19254	HBV-1537 CHZ-7 amino stab1	c ₅ g ₅ c ₅ g ₅ uaa cUGAUgagccgcuuagccGaa lagaggu B	6488
1791	UGUAGGC A UAAAUUG	6139	19258	HBV-1791 CHZ-7 amino stab1	c ₅ a ₅ a ₅ u ₅ uua cUGAUgagccgcuuagccGaa lcuaca B	6489
1831	UUUUCAC C UCUGCCU	6140	19260	HBV-1831 CHZ-7 amino stab1	a ₅ g ₅ g ₅ c ₅ aga cUGAUgagccgcuuagccGaa lugaana B	6490
1832	UUUCACC U CUGCCUA	6141	19261	HBV-1832 CHZ-7 amino stab1	u ₅ a ₅ g ₅ s ₅ cag cUGAUgagccgcuuagccGaa lgugaaa B	6491
1872	UUCAAGC C UCCAAGC	6142	19262	HBV-1872 CHZ-7 amino stab1	g ₅ c ₅ u ₅ s ₅ gga cUGAUgagccgcuuagccGaa lcuugaa B	6492
1873	UCAAGCC C CCAAGCU	6143	19263	HBV-1873 CHZ-7 amino stab1	a ₅ g ₅ c ₅ s ₅ ugg cUGAUgagccgcuuagccGaa lgcunga B	6493
1875	AAGCCUC C AAGCUGU	6144	19264	HBV-1875 CHZ-7 amino stab1	a ₅ c ₅ a ₅ g ₅ cuu cUGAUgagccgcuuagccGaa laggcuu B	6494
1876	AGCCUCC A AGCUGUG	6145	19266	HBV-1876 CHZ-7 amino stab1	c ₅ a ₅ c ₅ a ₅ gcu cUGAUgagccgcuuagccGaa laggcuu B	6495
1880	UCCAAGC U GUGCCUU	6146	19268	HBV-1880 CHZ-7 amino stab1	a ₅ a ₅ g ₅ g ₅ cac cUGAUgagccgcuuagccGaa lcuugaa B	6496
2382	GAAGAAC U CCCUCGC	6147	19269	HBV-2382 CHZ-7 amino stab1	g ₅ c ₅ g ₅ a ₅ g99 cUGAUgagccgcuuagccGaa luucuc B	6497
2384	AGAACUC C CUCGCCU	6148	19270	HBV-2384 CHZ-7 amino stab1	a ₅ g ₅ g ₅ c ₅ gag cUGAUgagccgcuuagccGaa lcuugaa B	6498
2385	GAACUCC C UCGCCUC	6149	19271	HBV-2385 CHZ-7 amino stab1	g ₅ a ₅ g ₅ g ₅ cga cUGAUgagccgcuuagccGaa lcuugaa B	6499
2422	GCGUCGC A GAAGAUC	6150	19272	HBV-2422 CHZ-7 amino stab1	g ₅ a ₅ u ₅ c ₅ uuc cUGAUgagccgcuuagccGaa lgcagc B	6500
2830	CAUAUUC U UGGGAAC	6151	19273	HBV-2830 CHZ-7 amino stab1	g ₅ u ₅ u ₅ c ₅ cca cUGAUgagccgcuuagccGaa laauug B	6501
315	GCCAAAUUC G CAGUC	6152	20079	HBV-315 GCI.Rz-5/10 stab2	g ₅ a ₅ c ₅ g uGAU ₅ gcauGcacuagc gcg gaauuugc B	6502
381	AUCGCUGGAU G UGUCU	6153	20080	HBV-381 GCI.Rz-5/10 stab2	a ₅ g ₅ a ₅ a uGAU ₅ gcauGcacuagc gcg aauuugc B	6503
476	UUGCCCGUUU G UCCUC	6154	20081	HBV-476 GCI.Rz-5/10 stab2	g ₅ a ₅ g ₅ a uGAU ₅ gcauGcacuagc gcg aauuugc B	6504
694	AGUGCCAUUU G UUCAG	6155	20082	HBV-694 GCI.Rz-5/10 stab2	c ₅ u ₅ g ₅ a uGAU ₅ gcauGcacuagc gcg aauuugc B	6505
1265	CUCCUCUGCC G AUCCA	6156	20083	HBV-1265 GCI.Rz-5/10 stab2	u ₅ g ₅ g ₅ u uGAU ₅ gcauGcacuagc gcg ggcagagag B	6506
1601	CUUCACCUU G CACGU	6157	20084	HBV-1601 GCI.Rz-5/10 stab2	a ₅ c ₅ g ₅ g uGAU ₅ gcauGcacuagc gcg agaggugaag B	6507

Table 43

1881	CCUCCAAGCU G UGCCU	6158	20085	HBV-1881 GCI.Rz-5/10 stab2	a ₉ s ₉ s ₉ a uGAU ₉ g gcauGcacuagc gcg agcuuggagg B	6508
1883	UCCAAGCUGU G CCUUG	6159	20086	HBV-1883 GCI.Rz-5/10 stab2	c ₉ s ₉ a ₉ g uGAU ₉ g gcauGcacuagc gcg acagcuugga B	6509
2388	GAACUCCUC G CCUCG	6160	20087	HBV-2388 GCI.Rz-5/10 stab2	c ₉ s ₉ a ₉ g uGAU ₉ g gcauGcacuagc gcg gaggagauuc B	6510
381	GCUGGAU G UGUUGC	6161	20091	HBV-381 Zin.Rz-7 amino stab2	g ₉ s ₉ a ₉ s ₉ aca Gccgaaag CGaGugaGGuCu auccagc B	6511
392	CUGCGG G UUUUAUC	6162	20092	HBV-392 Zin.Rz-7 amino stab2	g ₉ a ₉ u ₉ a ₉ aaa Gccgaaag CGaGugaGGuCu gccgcag B	6512
420	UCCUGCU G CUAUGCC	6163	20093	HBV-420 Zin.Rz-7 amino stab2	g ₉ s ₉ c ₉ a ₉ uag Gccgaaag CGaGugaGGuCu agcagga B	6513
648	UAUGGGA G UGGGCCU	6164	20094	HBV-648 Zin.Rz-7 amino stab2	a ₉ s ₉ s ₉ c ₉ cca Gccgaaag CGaGugaGGuCu ucccau B	6514
711	UCGUAGG G CUUUGCC	6165	20095	HBV-711 Zin.Rz-7 amino stab2	g ₉ s ₉ s ₉ a ₉ s ₉ aag Gccgaaag CGaGugaGGuCu ccuacga B	6515
1262	CUCCUCU G CCGAUCC	6166	20096	HBV-1262 Zin.Rz-7 amino stab2	g ₉ s ₉ a ₉ u ₉ s ₉ cg Gccgaaag CGaGugaGGuCu agaggag B	6516
1835	CACCUCU G CCUAAUC	6167	20097	HBV-1835 Zin.Rz-7 amino stab2	g ₉ a ₉ u ₉ u ₉ s ₉ agg Gccgaaag CGaGugaGGuCu agaggug B	6517
2388	CUCCUCU G CCUCGCA	6168	20098	HBV-2388 Zin.Rz-7 amino stab2	u ₉ s ₉ c ₉ s ₉ g ₉ agg Gccgaaag CGaGugaGGuCu gaggag B	6518
192	GACCCCU G CUCGUGU	6169	20099	HBV-192 Zin.Rz-7 amino stab2	a ₉ s ₉ a ₉ c ₉ g ₉ agg Gccgaaag CGaGugaGGuCu agggguc B	6519
198	UGCUCGU G UUAACAGG	6170	20100	HBV-198 Zin.Rz-7 amino stab2	c ₉ s ₉ u ₉ s ₉ uaa Gccgaaag CGaGugaGGuCu acgagca B	6520
315	AAAAUUC G CAGUCCC	6171	20101	HBV-315 Zin.Rz-7 amino stab2	g ₉ s ₉ s ₉ a ₉ s ₉ cug Gccgaaag CGaGugaGGuCu gaauuuu B	6521
383	GGAUGU G UGUGCG	6172	20102	HBV-383 Zin.Rz-6 amino stab2	c ₉ s ₉ c ₉ a ₉ s ₉ ga Gccgaaag CGaGugaGGuCu acaucc B	6522
383	UGGAUGU G UCUGCGG	6173	20103	HBV-383 Zin.Rz-7 amino stab2	c ₉ s ₉ g ₉ s ₉ c ₉ aga Gccgaaag CGaGugaGGuCu acaucca B	6523
387	GUGUCU G CGGCGU	6174	20104	HBV-387 Zin.Rz-6 amino stab2	a ₉ s ₉ g ₉ s ₉ c ₉ cg Gccgaaag CGaGugaGGuCu agacac B	6524
390	GUCUGCG G CGUUUAU	6175	20105	HBV-390 Zin.Rz-7 amino stab2	u ₉ a ₉ a ₉ s ₉ acg Gccgaaag CGaGugaGGuCu cgcagac B	6525
392	UGCGGC G UUUUAU	6176	20106	HBV-392 Zin.Rz-6 amino stab2	a ₉ u ₉ s ₉ a ₉ s ₉ aa Gccgaaag CGaGugaGGuCu gccgca B	6526
425	UGCJAU G CCUCAU	6177	20107	HBV-425 Zin.Rz-6 amino stab2	a ₉ u ₉ s ₉ a ₉ s ₉ gg Gccgaaag CGaGugaGGuCu auagca B	6527
425	CUGCUAU G CCUCAUC	6178	20108	HBV-425 Zin.Rz-7 amino stab2	g ₉ s ₉ u ₉ s ₉ agg Gccgaaag CGaGugaGGuCu auagcag B	6528
468	GUAUGUU G CCCGUUU	6179	20109	HBV-468 Zin.Rz-7 amino stab2	a ₉ a ₉ a ₉ s ₉ ggg Gccgaaag CGaGugaGGuCu aacauac B	6529
476	CCCGUUU G UCCUCUA	6180	20110	HBV-476 Zin.Rz-7 amino stab2	u ₉ a ₉ g ₉ a ₉ ggga Gccgaaag CGaGugaGGuCu aaacggg B	6530
648	AUGGGA G UGGGCC	6181	20111	HBV-648 Zin.Rz-6 amino stab2	g ₉ s ₉ c ₉ s ₉ c ₉ ca Gccgaaag CGaGugaGGuCu ucccau B	6531
694	GCCAUUU G UUCAGUG	6182	20112	HBV-694 Zin.Rz-7 amino stab2	c ₉ s ₉ c ₉ u ₉ gaa Gccgaaag CGaGugaGGuCu aaauaggc B	6532
699	UUGUJUA G UGGUJUCG	6183	20113	HBV-699 Zin.Rz-7 amino stab2	c ₉ s ₉ a ₉ s ₉ cca Gccgaaag CGaGugaGGuCu ugaacaa B	6533
1262	UCCUCU G CCGAUC	6184	20114	HBV-1262 Zin.Rz-6 amino stab2	g ₉ a ₉ u ₉ c ₉ gg Gccgaaag CGaGugaGGuCu agagga B	6534
1440	CCCGUCG G CGCUGAA	6185	20115	HBV-1440 Zin.Rz-7 amino stab2	u ₉ u ₉ c ₉ a ₉ ggg Gccgaaag CGaGugaGGuCu cgacggg B	6535
1526	CACGGG G CGCACC	6186	20116	HBV-1526 Zin.Rz-6 amino stab2	g ₉ g ₉ u ₉ g ₉ c ₉ g Gccgaaag CGaGugaGGuCu cccgug B	6536
1526	CCACGGG G CGCACCU	6187	20117	HBV-1526 Zin.Rz-7 amino stab2	a ₉ s ₉ s ₉ u ₉ gg Gccgaaag CGaGugaGGuCu cccgugg B	6537
1557	CCCGUCU G UGCCUUC	6188	20118	HBV-1557 Zin.Rz-7 amino stab2	g ₉ s ₉ a ₉ s ₉ g ₉ ca Gccgaaag CGaGugaGGuCu agacggg B	6538

Table 43

1559	CGUCUGU G CCUUCUC	6189	20119	HBV-1559 Zin.Rz-7 amino stab2	g _s a _s g _s a _s agg Gccgaaag GCGaGugaGGuCu acagacg B	6539
1590	GCACUUC G CUUCACC	6190	20120	HBV-1590 Zin.Rz-7 amino stab2	g _s g _s u _s g _s aag Gccgaaag GCGaGugaGGuCu gaagugc B	6540
1835	ACCUCU G CCUAAU	6191	20121	HBV-1835 Zin.Rz-6 amino stab2	a _s u _s u _s a _s g _s Gccgaaag GCGaGugaGGuCu agaggu B	6541
2311	ACCAAU G CCCUAU	6192	20122	HBV-2311 Zin.Rz-7 amino stab2	a _s u _s a _s g _s g _s g _s Gccgaaag GCGaGugaGGuCu auuuggu B	6542
2420	CCGCGUC G CAGAAGA	6193	20123	HBV-2420 Zin.Rz-7 amino stab2	u _s c _s u _s u _s cug Gccgaaag GCGaGugaGGuCu gacgagg B	6543
65	CCUGCUG G UGGCUCC	6194	20124	HBV-65 Zin.Rz-7 amino stab2	g _s g _s a _s g _s cca Gccgaaag GCGaGugaGGuCu cagcagg B	6544
192	ACCCCU G CUCGUG	6195	20125	HBV-192 Zin.Rz-6 amino stab2	c _s a _s c _s g _s ag Gccgaaag GCGaGugaGGuCu aggggu B	6545
198	GCUCGU G UIACAG	6196	20126	HBV-198 Zin.Rz-6 amino stab2	c _s u _s g _s u _s aa Gccgaaag GCGaGugaGGuCu acgagc B	6546
258	UAGACUC G UGGUGGA	6197	20127	HBV-258 Zin.Rz-7 amino stab2	u _s c _s c _s a _s cca Gccgaaag GCGaGugaGGuCu gagucua B	6547
261	ACUCGUG G UGGACUU	6198	20128	HBV-261 Zin.Rz-7 amino stab2	a _s a _s g _s u _s cca Gccgaaag GCGaGugaGGuCu cagcagu B	6548
315	AAAUUC G CAGUCC	6199	20129	HBV-315 Zin.Rz-6 amino stab2	g _s g _s a _s c _s ug Gccgaaag GCGaGugaGGuCu gaaauu B	6549
381	CUGGAU G UGUCUG	6200	20130	HBV-381 Zin.Rz-6 amino stab2	c _s a _s g _s a _s ca Gccgaaag GCGaGugaGGuCu auccag B	6550
387	UGUGUCU G CGGCGUU	6201	20131	HBV-387 Zin.Rz-7 amino stab2	a _s a _s c _s g _s cog Gccgaaag GCGaGugaGGuCu aaccag B	6551
390	UCUGCG G CGUUUU	6202	20132	HBV-390 Zin.Rz-6 amino stab2	a _s a _s a _s a _s c _s g Gccgaaag GCGaGugaGGuCu cgcaga B	6552
417	CAUCCU G CUGCUA	6203	20133	HBV-417 Zin.Rz-6 amino stab2	u _s a _s g _s c _s ag Gccgaaag GCGaGugaGGuCu aggaug B	6553
420	CCUGCU G CUAUGC	6204	20134	HBV-420 Zin.Rz-6 amino stab2	g _s c _s a _s u _s ag Gccgaaag GCGaGugaGGuCu agcagg B	6554
468	UAUGUU G CCCGUU	6205	20135	HBV-468 Zin.Rz-6 amino stab2	a _s a _s c _s g _s g _s g _s Gccgaaag GCGaGugaGGuCu aacaua B	6555
476	CCGUUU G UCCUCU	6206	20136	HBV-476 Zin.Rz-6 amino stab2	a _s g _s a _s g _s ga Gccgaaag GCGaGugaGGuCu aaacgg B	6556
677	GGCUCA G UUUACU	6207	20137	HBV-677 Zin.Rz-6 amino stab2	a _s g _s u _s a _s aa Gccgaaag GCGaGugaGGuCu ugagcc B	6557
677	UGGCUCA G UUUACUA	6208	20138	HBV-677 Zin.Rz-7 amino stab2	u _s a _s g _s u _s aaa Gccgaaag GCGaGugaGGuCu ugagcca B	6558
685	UUACUA G UGCCAU	6209	20139	HBV-685 Zin.Rz-6 amino stab2	a _s u _s g _s g _s ca Gccgaaag GCGaGugaGGuCu uaguaa B	6559
685	UUUACUA G UGCCAUU	6210	20140	HBV-685 Zin.Rz-7 amino stab2	a _s a _s u _s g _s g _s ca Gccgaaag GCGaGugaGGuCu uaguaaa B	6560
687	UACUAGU G CCAUUUG	6211	20141	HBV-687 Zin.Rz-7 amino stab2	c _s a _s a _s a _s ugg Gccgaaag GCGaGugaGGuCu acuagua B	6561
699	UGUUCA G UGGUUC	6212	20142	HBV-699 Zin.Rz-6 amino stab2	g _s a _s a _s c _s ca Gccgaaag GCGaGugaGGuCu ugaaca B	6562
702	UCAGUG G UUCGUA	6213	20143	HBV-702 Zin.Rz-6 amino stab2	u _s a _s c _s g _s aa Gccgaaag GCGaGugaGGuCu cacuga B	6563
702	UUCAGUG G UUCGUAG	6214	20144	HBV-702 Zin.Rz-7 amino stab2	c _s u _s a _s c _s g _s aa Gccgaaag GCGaGugaGGuCu cacugaa B	6564
711	CGUAGG G CUUUC	6215	20145	HBV-711 Zin.Rz-6 amino stab2	g _s g _s a _s a _s ag Gccgaaag GCGaGugaGGuCu ccuaog B	6565
1006	UUGUGG G UCUUUU	6216	20146	HBV-1006 Zin.Rz-6 amino stab2	a _s a _s a _s a _s ga Gccgaaag GCGaGugaGGuCu ccacaa B	6566
1103	UUUCUC G CCAACU	6217	20147	HBV-1103 Zin.Rz-6 amino stab2	a _s g _s u _s g _s gg Gccgaaag GCGaGugaGGuCu gagaaa B	6567
1103	CUUUCUC G CCAACUU	6218	20148	HBV-1103 Zin.Rz-7 amino stab2	a _s a _s g _s u _s ugg Gccgaaag GCGaGugaGGuCu gagaaag B	6568
1184	GCCAAAGU G UUUUCUG	6219	20149	HBV-1184 Zin.Rz-7 amino stab2	c _s a _s g _s c _s aaa Gccgaaag GCGaGugaGGuCu acuuggc B	6569

Table 43

1440	CCGUCG G CGCUGA	6220	20150	HBV-1440 Zin.Rz-6 amino stab2	U ₅ C ₅ A ₅ G ₅ C ₅ G GcogaagGCGaGugaGGuCu cgaagg B	6570
1442	GUCGGC G CUGAAU	6221	20151	HBV-1442 Zin.Rz-6 amino stab2	A ₅ U ₅ U ₅ C ₅ ag GcogaagGCGaGugaGGuCu gccgac B	6571
1442	CGUCGGC G CUGAAUC	6222	20152	HBV-1442 Zin.Rz-7 amino stab2	G ₅ A ₅ U ₅ U ₅ C ₅ ag GcogaagGCGaGugaGGuCu gccgacg B	6572
1553	CUCCCC G UCUGUG	6223	20153	HBV-1553 Zin.Rz-6 amino stab2	C ₅ A ₅ C ₅ A ₅ G GcogaagGCGaGugaGGuCu ggggag B	6573
1557	CCGUCU G UGCCUU	6224	20154	HBV-1557 Zin.Rz-6 amino stab2	A ₅ A ₅ G ₅ G ₅ C ₅ ca GcogaagGCGaGugaGGuCu agacgg B	6574
1559	GUCUGU G CCUUCU	6225	20155	HBV-1559 Zin.Rz-6 amino stab2	A ₅ G ₅ A ₅ A ₅ gg GcogaagGCGaGugaGGuCu acagac B	6575
1583	CCGUGU G CACUUC	6226	20156	HBV-1583 Zin.Rz-6 amino stab2	G ₅ A ₅ A ₅ G ₅ ug GcogaagGCGaGugaGGuCu acacgg B	6576
1590	CACUUC G CUUCAC	6227	20157	HBV-1590 Zin.Rz-6 amino stab2	G ₅ U ₅ G ₅ A ₅ ag GcogaagGCGaGugaGGuCu gaagug B	6577
1622	ACCACC G UGAACG	6228	20158	HBV-1622 Zin.Rz-6 amino stab2	C ₅ G ₅ U ₅ U ₅ ca GcogaagGCGaGugaGGuCu gguggu B	6578
1870	UGUUCAA G CCUCCAA	6229	20159	HBV-1870 Zin.Rz-7 amino stab2	U ₅ U ₅ G ₅ G ₅ agg GcogaagGCGaGugaGGuCu uugaaca B	6579
1881	CCAAGCU G UGCCUUG	6230	20160	HBV-1881 Zin.Rz-7 amino stab2	C ₅ A ₅ A ₅ G ₅ Gca GcogaagGCGaGugaGGuCu agcuugg B	6580
1883	AGCUGU G CCUUGG	6231	20161	HBV-1883 Zin.Rz-6 amino stab2	C ₅ C ₅ A ₅ A ₅ gg GcogaagGCGaGugaGGuCu acagcu B	6581
1883	AAGCUGU G CCUUGGG	6232	20162	HBV-1883 Zin.Rz-7 amino stab2	C ₅ C ₅ C ₅ A ₅ agg GcogaagGCGaGugaGGuCu acagcuu B	6582
2311	CCAAU G CCCCUA	6233	20163	HBV-2311 Zin.Rz-6 amino stab2	U ₅ A ₅ G ₅ G ₅ gg GcogaagGCGaGugaGGuCu auuugg B	6583
2347	ACUGUU G UUAGAC	6234	20164	HBV-2347 Zin.Rz-6 amino stab2	G ₅ U ₅ C ₅ U ₅ aa GcogaagGCGaGugaGGuCu aacagu B	6584
2364	AGGCAG G UCCCCU	6235	20165	HBV-2364 Zin.Rz-6 amino stab2	A ₅ G ₅ G ₅ G ₅ ga GcogaagGCGaGugaGGuCu cugccu B	6585
2364	GAGGCAG G UCCCCUA	6236	20166	HBV-2364 Zin.Rz-7 amino stab2	U ₅ A ₅ G ₅ G ₅ gg GcogaagGCGaGugaGGuCu cugccuc B	6586
2388	UCCCU G CCUCGC	6237	20167	HBV-2388 Zin.Rz-6 amino stab2	G ₅ C ₅ G ₅ A ₅ gg GcogaagGCGaGugaGGuCu gagggg B	6587
2393	CGCCUC G CAGACG	6238	20168	HBV-2393 Zin.Rz-6 amino stab2	C ₅ G ₅ U ₅ C ₅ ug GcogaagGCGaGugaGGuCu gagggc B	6588
2417	CGCCGC G UGCAG	6239	20169	HBV-2417 Zin.Rz-6 amino stab2	C ₅ U ₅ G ₅ C ₅ ga GcogaagGCGaGugaGGuCu ggggag B	6589
2420	CGCGUC G CAGAAG	6240	20170	HBV-2420 Zin.Rz-6 amino stab2	C ₅ U ₅ U ₅ C ₅ ug GcogaagGCGaGugaGGuCu gaagcg B	6590
2474	CAUAAG G UGGGAA	6241	20171	HBV-2474 Zin.Rz-6 amino stab2	U ₅ U ₅ C ₅ C ₅ ca GcogaagGCGaGugaGGuCu cuuaug B	6591
381	GCUGGAU G UGUCUGC	6242	20172	HBV-381 Amb.Rz-7 stab2	G ₅ C ₅ A ₅ G ₅ aca gga L ucCCUUCaagga L ucCGGG auccagc B	6592
648	UAUGGA G UGGGCCU	6243	20173	HBV-648 Amb.Rz-7 stab2	A ₅ G ₅ G ₅ C ₅ cca gga L ucCCUUCaagga L ucCGGG ucccau B	6593
198	UGCUCGU G UIACAGG	6244	20174	HBV-198 Amb.Rz-7 stab2	C ₅ C ₅ U ₅ G ₅ uaa gga L ucCCUUCaagga L ucCGGG acgagca B	6594
377	UAUCGCU G GAUGUGU	6245	20175	HBV-377 Amb.Rz-7 stab2	A ₅ C ₅ A ₅ C ₅ auc gga L ucCCUUCaagga L ucCGGG agcgaua B	6595
378	AUCGCU G AUGUGUC	6246	20176	HBV-378 Amb.Rz-7 stab2	G ₅ A ₅ C ₅ A ₅ cau gga L ucCCUUCaagga L ucCGGG cagcgau B	6596
383	UGGAUGU G UCUGCGG	6247	20177	HBV-383 Amb.Rz-7 stab2	C ₅ C ₅ G ₅ C ₅ aga gga L ucCCUUCaagga L ucCGGG acaucca B	6597
383	GGAUGU G UCUGCG	6248	20178	HBV-383 Amb.Rz-6 stab2	C ₅ G ₅ C ₅ A ₅ ga gga L ucCCUUCaagga L ucCGGG acaucc B	6598
648	AUGGA G UGGGCC	6249	20179	HBV-648 Amb.Rz-6 stab2	G ₅ G ₅ C ₅ C ₅ ca gga L ucCCUUCaagga L ucCGGG ucccau B	6599
650	UGGGAGU G GGCCUCA	6250	20180	HBV-650 Amb.Rz-7 stab2	U ₅ G ₅ A ₅ G ₅ gcc gga L ucCCUUCaagga L ucCGGG acucca B	6600

Table 43

650	GGGAGU G GGCCUC	6251	20181	HBV-650 Amb.Rz-6 stab2	g ₅ a ₅ g ₅ cc gga L ucCCUUUcaagga L ucCGGG acucc B	6601
694	GCCAUUU G UUCAGUG	6252	20182	HBV-694 Amb.Rz-7 stab2	c ₅ a ₅ c ₅ u ₅ gaa gga L ucCCUUUcaagga L ucCGGG aauggc B	6602
699	UUGUUA G UGUUUG	6253	20183	HBV-699 Amb.Rz-7 stab2	c ₅ g ₅ a ₅ c ₅ cca gga L ucCCUUUcaagga L ucCGGG ugaaca B	6603
701	GUUCAGU G GUUCGUA	6254	20184	HBV-701 Amb.Rz-7 stab2	u ₅ a ₅ c ₅ g ₅ aac gga L ucCCUUUcaagga L ucCGGG acugaac B	6604
710	UUCGUAG G GCUUUC	6255	20185	HBV-710 Amb.Rz-7 stab2	g ₅ g ₅ a ₅ g ₅ agc gga L ucCCUUUcaagga L ucCGGG cuacga B	6605
1525	CCACGG G GCGCAC	6256	20186	HBV-1525 Amb.Rz-6 stab2	g ₅ u ₅ g ₅ c ₅ g ₅ c gga L ucCCUUUcaagga L ucCGGG ccgugg B	6606
1624	CACCGU G AACGCC	6257	20187	HBV-1624 Amb.Rz-6 stab2	g ₅ g ₅ c ₅ g ₅ uu gga L ucCCUUUcaagga L ucCGGG acggug B	6607
2069	CACUCA G GCAAGC	6258	20188	HBV-2069 Amb.Rz-6 stab2	g ₅ c ₅ u ₅ g ₅ c gga L ucCCUUUcaagga L ucCGGG ugagug B	6608
2375	CCUAGAA G AAGAACU	6259	20189	HBV-2375 Amb.Rz-7 stab2	a ₅ g ₅ u ₅ u ₅ cuu gga L ucCCUUUcaagga L ucCGGG uucuaag B	6609
2476	AUAAGGU G GGAAACU	6260	20190	HBV-2476 Amb.Rz-7 stab2	a ₅ g ₅ u ₅ u ₅ ucc gga L ucCCUUUcaagga L ucCGGG accuuu B	6610
65	CCUGCUG G UGCUCC	6261	20191	HBV-65 Amb.Rz-7 stab2	g ₅ g ₅ a ₅ g ₅ cca gga L ucCCUUUcaagga L ucCGGG cagcagg B	6611
67	GCUGGU G GCUCCA	6262	20192	HBV-67 Amb.Rz-6 stab2	u ₅ g ₅ g ₅ a ₅ g ₅ c gga L ucCCUUUcaagga L ucCGGG accagc B	6612
198	GCUCGU G UUAACAG	6263	20193	HBV-198 Amb.Rz-6 stab2	c ₅ u ₅ g ₅ u ₅ aa gga L ucCCUUUcaagga L ucCGGG acgagc B	6613
260	GACUCGU G GUGGACU	6264	20194	HBV-260 Amb.Rz-7 stab2	a ₅ g ₅ u ₅ c ₅ cac gga L ucCCUUUcaagga L ucCGGG acgaguc B	6614
263	UCGUGGU G GACUUCU	6265	20195	HBV-263 Amb.Rz-7 stab2	a ₅ g ₅ a ₅ g ₅ guc gga L ucCCUUUcaagga L ucCGGG accacga B	6615
377	AUCGCU G GAUGUG	6266	20196	HBV-377 Amb.Rz-6 stab2	c ₅ a ₅ c ₅ a ₅ uc gga L ucCCUUUcaagga L ucCGGG agcgau B	6616
378	UCGCU G AUGUGU	6267	20197	HBV-378 Amb.Rz-6 stab2	a ₅ c ₅ a ₅ c ₅ au gga L ucCCUUUcaagga L ucCGGG cagcga B	6617
476	CCGUUU G UCCUCU	6268	20198	HBV-476 Amb.Rz-6 stab2	a ₅ g ₅ a ₅ g ₅ ga gga L ucCCUUUcaagga L ucCGGG aaacgg B	6618
651	GGGAGUG G GCCUCAG	6269	20199	HBV-651 Amb.Rz-7 stab2	c ₅ u ₅ g ₅ a ₅ g ₅ c gga L ucCCUUUcaagga L ucCGGG cacucc B	6619
677	UGGCUCA G UUUACUA	6270	20200	HBV-677 Amb.Rz-7 stab2	u ₅ a ₅ g ₅ u ₅ aaa gga L ucCCUUUcaagga L ucCGGG ugagcca B	6620
685	UUUACUA G UGCCAUU	6271	20201	HBV-685 Amb.Rz-7 stab2	a ₅ a ₅ u ₅ g ₅ gca gga L ucCCUUUcaagga L ucCGGG uaguaaa B	6621
702	UUCAGUG G UUCGUAG	6272	20202	HBV-702 Amb.Rz-7 stab2	c ₅ u ₅ a ₅ c ₅ gaa gga L ucCCUUUcaagga L ucCGGG cacuga B	6622
709	GUUCGUA G GGCUUUC	6273	20203	HBV-709 Amb.Rz-7 stab2	g ₅ a ₅ a ₅ g ₅ gcc gga L ucCCUUUcaagga L ucCGGG uacgaac B	6623
710	UCGUAG G GCUUUC	6274	20204	HBV-710 Amb.Rz-6 stab2	g ₅ a ₅ a ₅ g ₅ c gga L ucCCUUUcaagga L ucCGGG cuacga B	6624
747	UAUGGAU G AUGUGGU	6275	20205	HBV-747 Amb.Rz-7 stab2	a ₅ c ₅ c ₅ a ₅ cau gga L ucCCUUUcaagga L ucCGGG auccaua B	6625
1557	CCGUCU G UGCCUU	6276	20206	HBV-1557 Amb.Rz-6 stab2	a ₅ a ₅ g ₅ g ₅ ca gga L ucCCUUUcaagga L ucCGGG agacgg B	6626
1881	CCAAGCU G UGCCUUG	6277	20207	HBV-1881 Amb.Rz-7 stab2	c ₅ a ₅ g ₅ g ₅ gca gga L ucCCUUUcaagga L ucCGGG agcuugg B	6627
2347	ACUGUU G UUAGAC	6278	20208	HBV-2347 Amb.Rz-6 stab2	g ₅ u ₅ c ₅ u ₅ aa gga L ucCCUUUcaagga L ucCGGG aacagu B	6628
2375	CUAGAA G AAGAAC	6279	20209	HBV-2375 Amb.Rz-6 stab2	g ₅ u ₅ u ₅ c ₅ uu gga L ucCCUUUcaagga L ucCGGG uucuaag B	6629
2378	GAAGAA G AACUCC	6280	20210	HBV-2378 Amb.Rz-6 stab2	g ₅ g ₅ a ₅ g ₅ uu gga L ucCCUUUcaagga L ucCGGG uucuu B	6630
2423	CGUCGCA G AAGAUCU	6281	20211	HBV-2423 Amb.Rz-7 stab2	a ₅ g ₅ a ₅ u ₅ cuu gga L ucCCUUUcaagga L ucCGGG ugacag B	6631

Table 43

2426	GCAGAA G AUCUCA	6282	20212	HBV-2426 Amb.Rz-6 stab2	u _s s _s a _s s _s au gga L ucCCUU ^{Caagga} L ucCGGG uucugc B	6632
2426	CGCAGAA G AUCUCA	6283	20213	HBV-2426 Amb.Rz-7 stab2	u _s u _s g _s a _s s _s g _s au gga L ucCCUU ^{Caagga} L ucCGGG uucugc B	6633
2476	UAAGGU G GGAAC	6284	20214	HBV-2476 Amb.Rz-6 stab2	g _s u _s u _s u _s cc gga L ucCCUU ^{Caagga} L ucCGGG accuua B	6634
2477	UAAGGUG G GAAACUU	6285	20215	HBV-2477 Amb.Rz-7 stab2	a _s s _s g _s u _s uuc gga L ucCCUU ^{Caagga} L ucCGGG caccuua B	6635
2477	AAGGUG G GAAACU	6286	20216	HBV-2477 Amb.Rz-6 stab2	a _s g _s u _s u _s uc gga L ucCCUU ^{Caagga} L ucCGGG caccuu B	6636
1607	UGCACGU C GCAUGGA	6287	20697	HBV-1607 Rz-7 allyl stab1 (7/4)	u _s c _s c _s a _s u _s g _s cUGAuGaggccguuagccGaa Acgugca B	6637
1887	GUGCCU U GGGUGG	6288	20698	HBV-1887 Rz-6 allyl stab1 (6/4)	c _s c _s a _s c _s cc cUGAuGaggccguuagccGaa Aggcac B	6638
1607	GCACGU C GCAUGG	6289	20699	HBV-1607 Rz-6 allyl stab1 (6/3)	c _s c _s a _s u _s g _s cc cUGAuGaggccguuagccGaa Acgugc B	6639
1607	UGCACGU C GCAUGGA	6290	20700	HBV-1607 Rz-7 allyl stab1 (7/3)	u _s c _s c _s a _s u _s g _s cc cUGAuGaggccguuagccGaa Acgugca B	6640
1887	GUGCCU U GGGUGG	6291	20701	HBV-1887 Rz-6 allyl stab1 (6/3)	c _s c _s a _s c _s cc cUGAuGaggccguuagccGaa Aggcac B	6641
1887	UGUGCCU U GGGUGGC	6292	20702	HBV-1887 Rz-7 allyl stab1 (7/3)	g _s c _s c _s a _s ccc cUGAuGaggccguuagccGaa Aggcaca B	6642

UPPER CASE = RIBO

lower case = 2'-O-methyl

s = phosphorothioate linkage

B = inverted deoxybasic residue

U = 2'-deoxy-2'-C-allyl Uridine

U = 2'-deoxy-2'-amino Uridine

C = 2'-deoxy-2'-amino Cytidine

Table 44

Table 44: Group Designation and Dosage levels for HBV transgenic mouse study

Group	Compound	Dose	Number of Mice	Duration of Treatment
1	RPI.18341 (site 273)	100 mg/kg/day*	10F	14 days
2	RPI.18371 (site 1833)	100 mg/kg/day*	10F	14 days
3	RPI.18418 (site 1873)	100 mg/kg/day*	10F	14 days
4	RPI.18372 (site 1874)	100 mg/kg/day*	10F	14 days
5	Saline control	100 mg/kg/day*	10F	14 days
6	Untreated		10F	0 days

*administered via sc infusion using Alzet mini-osmotic pumps

Table 45

TABLE 45. NUCLEOSIDES USED FOR CHEMICAL SYNTHESIS OF MODIFIED NUCLEOTIDE TRIPHOSPHATES

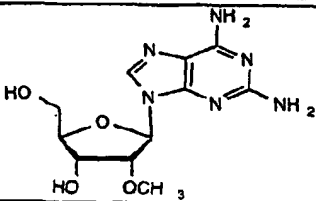
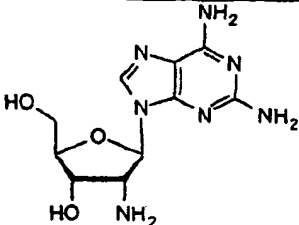
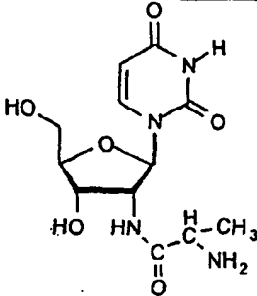
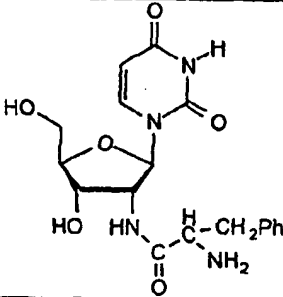
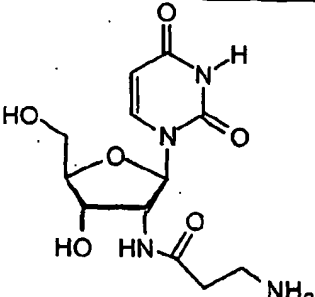
	NUCLEOSIDES	Abbreviation	CHEMICAL STRUCTURE
1	2'-O-methyl-2,6-diaminopurine riboside	2'-O-Me-DAP	
2	2'-deoxy-2' amino-2,6-diaminopurine riboside	2'-NH ₂ -DAP	
3	2'-(N-alanyl)amino-2'-deoxy-uridine	ala-2'- NH ₂ U	
4	2'-(N-phenylalanyl)amino-2'-deoxy-uridine	phe-2'- NH ₂ -U	
5	2'-(N-β-alanyl) amino-2'-deoxy uridine	2'-β-Ala-NH ₂ -U	

Table 45

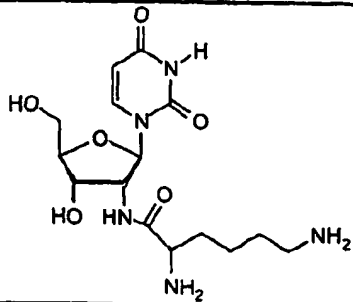
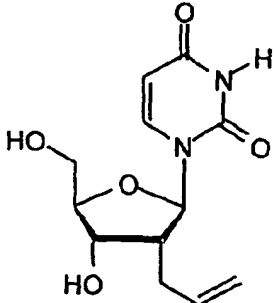
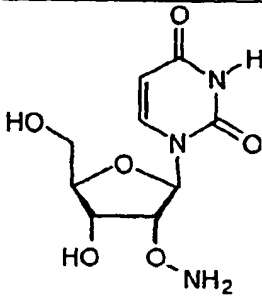
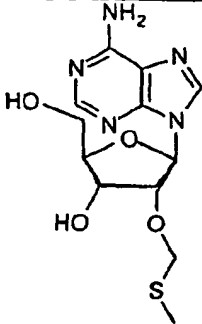
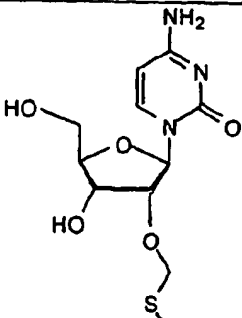
6	2'-Deoxy-2'-(lysiyl) amino uridine	2'-L-lys-NH ₂ U	
7	2'-C-allyl uridine	2'-C-allyl-U	
8	2'-O-amino-uridine	2'-O-NH ₂ -U	
9	2'-O-methylthiomethyl adenosine	2'-O-MTM-A	
10	2'-O-methylthiomethyl cytidine	2'-O-MTM-C	

Table 45

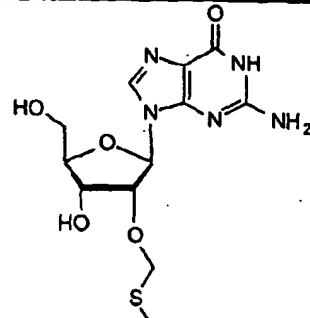
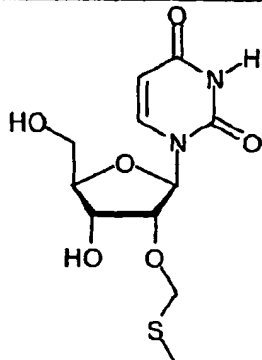
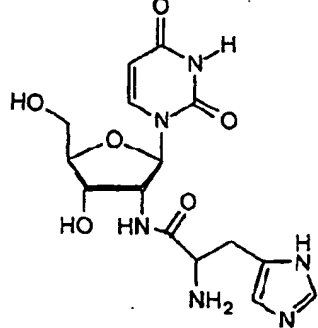
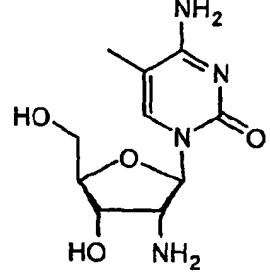
11	2'-O-methylthiomethyl guanosine	2'-O-MTM-G	
12	2'-O-methylthiomethyl- uridine	2'-O-MTM-U	
13	2'-(N-histidyl) amino uridine	2'-his-NH ₂ -U	
14	2'-Deoxy-2'-amino-5- methyl cytidine	5-Me-2'-NH ₂ -C	

Table 45

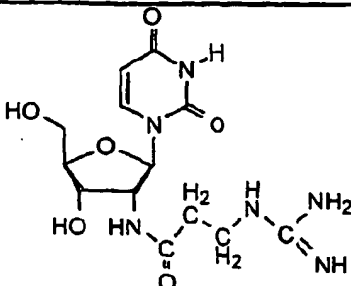
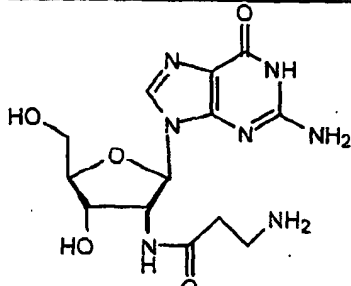
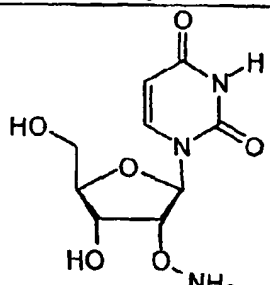
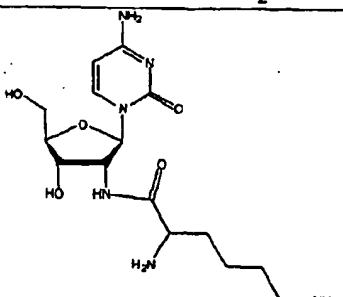
15	2'-(N-β-carboxamidine-β-alanyl)amino-2'-deoxy-uridine	β-ala-CA-NH ₂ -U	
16	2'-(N-β-alanyl) guanosine	β-Ala-NH ₂ -G	
17	2'-O-Amino-Uridine	2'-O-NH ₂ -U	
18	2'-(N-lysyl)amino -2'-deoxy-cytidine	2'-NH ₂ -lys-C	

Table 45

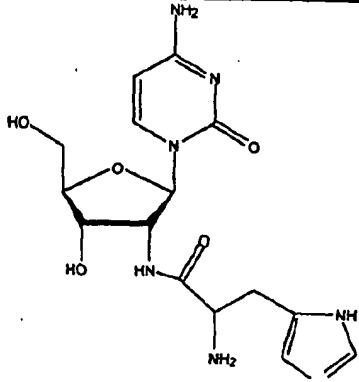
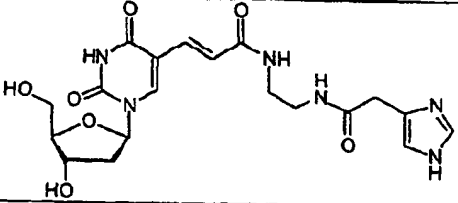
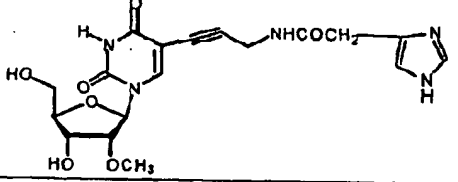
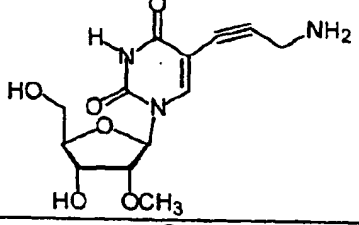
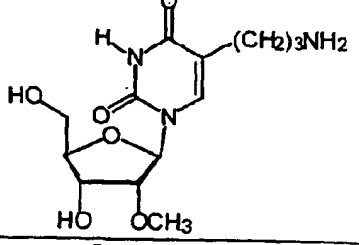
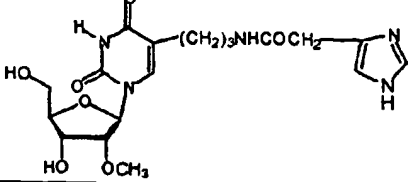
19	2'-Deoxy histidine) Cytidine	-2'-(L- amino	2'- NH ₂ -his-C	
20	5-Imidazoleacetic acid 2'-deoxy uridine		5-IAA-U	
21	5-[3-(N-4- imidazoleacetyl)amino propynyl]-2'-O-methyl uridine		5-IAA- propynylamino- 2'-OMe U	
22	5-(3-aminopropynyl)- 2'-O-methyl uridine		5-aminopropynyl- 2'-OMe U	
23	5-(3-aminopropyl)-2'- O-methyl uridine		5-aminopropyl- 2'-OMe U	
24	5-[3-(N-4- imidazoleacetyl)amino propyl]-2'-O-methyl Uridine		5-IAA- propylamino-2'- OMe U	

Table 45

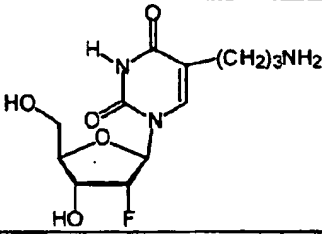
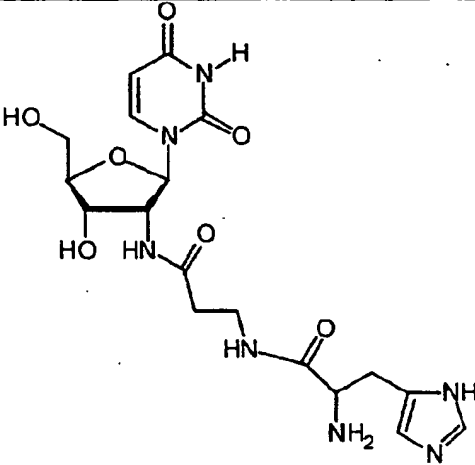
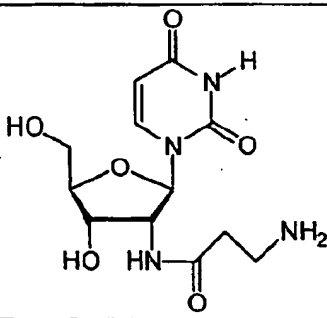
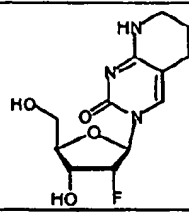
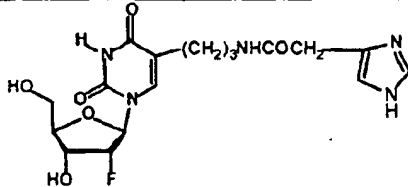
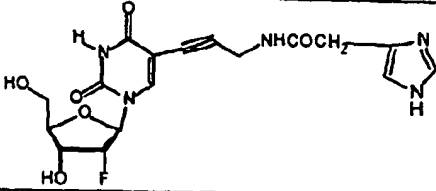
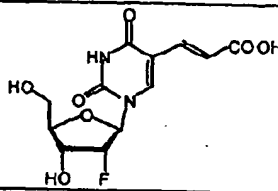
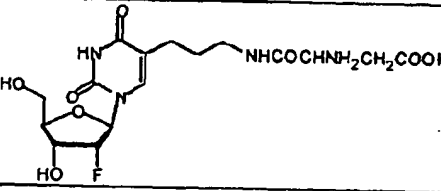
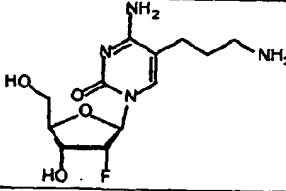
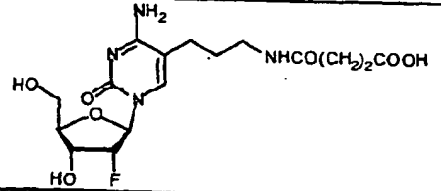
25	5-(3-aminopropyl)-2'-deoxy-2-fluoro uridine	5-aminopropyl-2'-F dU	
26	2'-Deoxy-2'-(β-alanyl-L-histidyl)amino Uridine	2'-amino-β-ALA-HIS dU	
27	2'-deoxy-2'-β-alaninamido-uridine	2'-β-ALA dU	
28	3-(2'-deoxy-2'-fluoro-β-D-ribofuranosyl)piperazino[2,3-D]pyrimidine-2-one	2'-F piperazino-pyrimidinone	
29	5-[3-(N-4-imidazoleacetyl)amino propyl]-2'-deoxy-2'-fluoro Uridine	5-IAA-propylamino-2'-F dU	

Table 45

30	5-[3-(N-4-imidazoleacetyl)amino propynyl]-2'-deoxy-2'-fluoro uridine	5-IAA-propynylamino-2'-F dU	
31	5-E-(2-carboxyvinyl-2'-deoxy-2'-fluoro uridine	5-carboxyvinyl-2'-F dU	
32	5-[3-(N-4-aspartyl)aminopropynyl]-2'-fluoro uridine	5-ASP-aminopropyl-2'-F-dU	
33	5-(3-aminopropyl)-2'-deoxy-2'-fluoro cytidine	5-aminopropyl-2'-F dC	
34	5-[3-(N-4-succinyl)aminopropyl]-2'-deoxy-2'-fluoro cytidine	5-succinylamino-propyl-2'-F dC	

605
Table 46

Table 46: PHOSPHORYLATION OF URIDINE IN THE PRESENCE OF DMAP

0 equiv. DMAP		0.2 equiv. DMAP		0.5 equiv. DMAP		1.0 equiv. DMAP	
Time (min)	Product %	Time (min)	Product %	Time (min)	Product %	Time (min)	Product %
0	1	0	0	0	0	0	0
40	7	10	8	20	27	30	74
80	10	50	24	60	46	70	77
120	12	90	33	100	57	110	84
160	14	130	39	140	63	150	83
200	17	170	43	180	63	190	84
240	19	210	47	220	64	230	77
320	20	250	48	260	68	270	79
1130	48	290	49	300	64	310	77
1200	46	1140	68	1150	76	1160	72
		1210	69	1220	76	1230	74

Table 47

Table 47: Detailed Description of the NTP Incorporation Reaction Conditions

Condition No.	TRIS-HCL (mM)	MgCl ₂ (mM)	DTT (mM)	Spermidine (mM)	Triton X-100 (%)	METHANOL (%)	LiCl (mM)	PEG (%)	Temp(°C)
1	40 (pH 8.0)	20	10	5	0.01	10	1	-	25
2	40 (pH 8.0)	20	10	5	0.01	10	1	4	25
3	40 (pH 8.1)	12	5	1	0.002	-	-	4	25
4	40 (pH 8.1)	12	5	1	0.002	10	-	4	25
5	40 (pH 8.1)	12	5	1	0.002	-	1	4	25
6	40 (pH 8.1)	12	5	1	0.002	10	1	4	25
7	40 (pH 8.0)	20	10	5	0.01	10	1	-	37
8	40 (pH 8.0)	20	10	5	0.01	10	1	4	37
9	40 (pH 8.1)	12	5	1	0.002	-	-	4	37
10	40 (pH 8.1)	12	5	1	0.002	10	-	4	37
11	40 (pH 8.1)	12	5	1	0.002	-	1	4	37
12	40 (pH 8.1)	12	5	1	0.002	10	1	4	37

Table 48

Table 48: INCORPORATION OF MODIFIED NUCLEOTIDE TRIPHOSPHATES

Modification	COND# 1	COND# 2	COND# 3	COND# 4	COND# 5	COND# 6	COND# 7	COND# 8	COND# 9	COND# 10	COND# 11	COND# 12
2'-NH ₂ -ATP	1	2	3	5	2	4	1	2	10	11	5	9
2'-NH ₂ -CTP	11	37	45	64	25	70	26	54	292	264	109	244
2'-NH ₂ -GTP	4	7	6	14	5	17	3	16	10	21	9	16
2'-NH ₂ -UTP	14	45	4	100	85	82	48	88	20	418	429	440
2'-dATP	9	3	19	23	9	24	6	3	84	70	28	51
2'-dCTP	1	10	43	46	35	47	27	127	204	212	230	235
2'-dGTP	6	10	9	15	9	12	8	34	38	122	31	46
2'-dTTP	9	9	14	18	13	18	8	15	116	114	59	130
2'-O-Me-ATP	0	0	0	0	0	0	1	1	2	2	2	2
2'-O-Me-CTP	no data compared to ribo; incorporates at low level											
2'-O-Me-GTP	4	3	4	4	4	4	2	4	4	5	4	5
2'-O-Me-UTP	55	52	39	38	41	48	55	71	93	103	81	77
2'-O-Me-DAP	4	4	3	4	4	5	4	3	4	5	5	5
2'-NH ₂ -DAP	0	0	1	1	1	1	1	0	0	0	0	0
ala-2'-NH ₂ -UTP	2	2	2	2	3	4	14	18	15	20	13	14
phe-2'-NH ₂ -UTP	8	12	7	7	8	8	4	10	6	6	10	6
2'-β NH ₂ -ala-UTP	65	48	25	17	21	21	220	223	265	300	275	248
2'-F-C5-carboxyvinyl UTP									100			
2'-F-C5-aspartyl-									100			

Table 49

**Table 49: INCORPORATION OF MODIFIED NUCLEOTIDE TRIPHOSPHATES
USING WILD TYPE BACTERIOPHAGE T7 POLYMERASE**

Modification	label	% ribo control
2'-NH ₂ -GTP	ATP	4%
2'-dGTP	ATP	3%
2'-O-Me-GTP	ATP	3%
2'-F-GTP	ATP	4%
2'-O-MTM-GTP	ATP	3%
2'-NH ₂ -UTP	ATP	39%
2'-dTTP	ATP	5%
2'-O-Me-UTP	ATP	3%
ala-2'-NH ₂ -UTP	ATP	2%
phe-2'-NH ₂ -UTP	ATP	1%
2'-β-ala-NH ₂ -UTP	ATP	3%
2'-C-allyl-UTP	ATP	2%
2'-O-NH ₂ -UTP	ATP	1%
2'-O-MTM-UTP	ATP	64%
2'-NH ₂ -ATP	GTP	1%
2'-O-MTM-ATP	GTP	1%
2'-NH ₂ -CTP	GTP	59%
2'-dCTP	GTP	40%
2'-F-CTP	GTP	100%
2'-F-UTP	GTP	100%
2'-F-TTP	GTP	0%
2'-F-C5-carboxyvinyl UTP	GTP	100%
2'-F-C5-aspartyl-aminopropyl UTP	GTP	100%
2'-F-C5-propylamine CTP	GTP	100%
2'-O-Me CTP	GTP	0%
2'-O-Me UTP	GTP	0%
2'-O-Me 5-3-aminopropyl UTP	GTP	0%
2'-O-Me 5-3-aminopropyl UTP	GTP	0%

Table 50

Table 50 a: Incorporation of 2'-his-UTP and Modified CTP's

modification	2'-his-UTP	rUTP
CTP	16.1	100
2'-amino-CTP	9.5*	232.7
2'-deoxy-CTP	9.6*	130.1
2'-OMe-CTP	1.9	6.2
2'-MTM-CTP	5.9	5.1
control	1.2	

Table 50 b: Incorporation of 2'-his-UTP, 2-amino CTP, and Modified ATP's

modification	2'-his-UTP and 2'-amino-CTP	rUTP and rCTP
ATP	15.7	100
2'-amino-ATP	2.4	28.9
2'-deoxy-ATP	2.3	146.3
2'-OMe-ATP	2.7	15
2'-F-ATP	4	222.6
2'-MTM-ATP	4.7	15.3
2'-OMe-DAP	1.9	5.7
2'-amino-DAP	8.9*	9.6

Numbers shown are a percentage of incorporation compared to the all-RNA control

* -Bold number indicates best observed rate of modified nucleotide triphosphate incorporation

Table 51

Table 51: INCORPORATION OF 2'-his-UTP, 2'-NH₂-CTP, 2'-NH₂-DAP, and rGTP USING VARIOUS REACTION CONDITIONS

Conditions	compared to all rNTP
7	8.7*
8	7*
9	2.3
10	2.7
11	1.6
12	2.5

Numbers shown are a percentage of incorporation compared to the all-RNA control

* Two highest levels of incorporation contained both methanol and LiCl

Table 52

Table 52: Selection of Oligonucleotides with Ribozyme Activity

pool	Generation	time	substrate remaining(%)	time	Substrate remaining (%)
N60	0	4 hr	100.00	24 hr	100.98
N60	14	4 hr	99.67	24 hr	97.51
N60	15	4 hr	98.76	24 hr	96.76
N60	16	4 hr	97.09	24 hr	96.60
N60	17	4 hr	79.50	24 hr	64.01
N40	0	4 hr	99.89	24 hr	99.78
N40	10	4 hr	99.74	24 hr	99.42
N40	11	4 hr	97.18	24 hr	90.38
N40	12	4 hr	61.64	24 hr	44.54
N40	13	4 hr	54.28	24 hr	36.46
N20	0	4 hr	99.18	24 hr	100.00
N20	11	4 hr	100.00	24 hr	100.00
N20	12	4 hr	99.51	24 hr	100.00
N20	13	4 hr	90.63	24 hr	84.89
N20	14	4 hr	91.16	24 hr	85.92
N60B	0	4 hr	100.00	24 hr	100.00
N60B	1	4 hr	100.00	24 hr	100.00
N60B	2	4 hr	100.00	24 hr	100.00
N60B	3	4 hr	100.00	24 hr	100.00
N60B	4	4 hr	99.24	24 hr	100.00
N60B	5	4 hr	97.81	24 hr	96.65
N60B	6	4 hr	89.95	24 hr	77.14

Table 53

Table 53: Kinetic Activity of Combinatorial Libraries

Pool	Generation	k_{obs} (min^{-1})
N60	17	0.0372
	18	0.0953
	19	0.0827
N40	12	0.0474
	13	0.037
	14	0.065
N20	15	0.0254
	13	0.0359
	14	0.0597
N60B	15	0.0549
	16	0.0477
	6	0.0209
	7	0.0715
	8	0.0379

Table 54

Table 54: Kinetic Activity of Clones within N60 and N40 Combinatorial Libraries

clone	library	activity(min ⁻¹)	k _{rel}
G18	N60	0.00226	1.00
0-2	N60	0.0389	17.21
0-3	N60	0.000609	0.27
0-5	N60	0.000673	0.30
0-7	N60	0.00104	0.46
0-8	N60	0.000739	0.33
0-11	N60	0.0106	4.69
0-12	N60	0.00224	0.99
0-13	N60	0.0255	11.28
0-14	N60	0.000878	0.39
0-15	N60	0.0000686	0.03
0-21	N60	0.0109	4.82
0-22	N60	0.000835	0.37
0-24	N60	0.000658	0.29
0-28	N40	0.000741	0.33
0-35	N40	0.00658	2.91
3-1	N40	0.0264	11.68
3-3	N40	0.000451	0.20
3-7	N40	0.000854	0.38
3-15	N40	0.000832	0.37

Table 55

Table 55: Effect of Magnesium Concentration of the Cleavage Rate of N20

[Mg⁺⁺]	k_{obs}(min⁻¹)
25	0.0259
20	0.0223
15	0.0182
10	0.0208
5	0.0121
2	0.00319
2	0.00226

Table 56

Enzymatic Nucleic Acid Motifs Targeting HCV

Seq ID	Alias	Sequence	Rz Seq ID
1	HCV.R1A-6 Amb.Rz-10/5	ggagugucgc GagggaacucC CU UCAAGGACAUUGUCCGGG ccacB	39
2	HCV.R1A-56 Amb.Rz-10/5	acgcunucug GagggaacucC CU UCAAGGACAUUGUCCGGG gugaa B	40
3	HCV.R1A-75 Amb.Rz-10/5	auacuaacgc GagggaacucC CU UCAAGGACAUUGUCCGGG auggc B	41
4	HCV.R1A-76 Amb.Rz-10/5	cauacuaacg GagggaacucC CU UCAAGGACAUUGUCCGGG caugg B	42
5	HCV.R1A-95 Amb.Rz-10/5	cuggagucug GagggaacucC CU UCAAGGACAUUGUCCGGG acgac B	43
6	HCV.R1A-138 Amb.Rz-10/5	accgguccg GagggaacucC CU UCAAGGACAUUGUCCGGG agacc B	44
7	HCV.R1A-146 Amb.Rz-10/5	guuacucac GagggaacucC CU UCAAGGACAUUGUCCGGG gguuc B	45
8	HCV.R1A-158 Amb.Rz-10/5	cuggcaauuc GagggaacucC CU UCAAGGACAUUGUCCGGG ggugu B	46
9	HCV.R1A-164 Amb.Rz-10/5	gucgucucg GagggaacucC CU UCAAGGACAUUGUCCGGG aauc B	47
10	HCV.R1A-176 Amb.Rz-10/5	agaaagacc GagggaacucC CU UCAAGGACAUUGUCCGGG ggucg B	48
11	HCV.R1A-177 Amb.Rz-10/5	aagaaagac GagggaacucC CU UCAAGGACAUUGUCCGGG cgguc B	49
12	HCV.R1A-209 Amb.Rz-10/5	cccaauuc GagggaacucC CU UCAAGGACAUUGUCCGGG agga B	50
13	HCV.R1A-237 Amb.Rz-10/5	acucggucag GagggaacucC CU UCAAGGACAUUGUCCGGG agucB	51
14	HCV.R1A-254 Amb.Rz-10/5	uuucgcacc GagggaacucC CU UCAAGGACAUUGUCCGGG aacac B	52
15	HCV.R1A-255 Amb.Rz-10/5	cuuucggac GagggaacucC CU UCAAGGACAUUGUCCGGG caaca B	53
16	HCV.R1A-259 Amb.Rz-10/5	agggcuucg GagggaacucC CU UCAAGGACAUUGUCCGGG gacc B	54
17	HCV.R1A-266 Amb.Rz-10/5	uaccacaag GagggaacucC CU UCAAGGACAUUGUCCGGG cuuc B	55
18	HCV.R1A-273 Amb.Rz-10/5	caggcaguac GagggaacucC CU UCAAGGACAUUGUCCGGG acaag B	56
19	HCV.R1A-288 Amb.Rz-10/5	ucgcaagcac GagggaacucC CU UCAAGGACAUUGUCCGGG cuac B	57
20	HCV.R1A-291 Amb.Rz-10/5	cacucgcaag GagggaacucC CU UCAAGGACAUUGUCCGGG acccu B	58
21	HCV.R1A-7 Amb.Rz-10/5	uggagucug GagggaacucC CU UCAAGGACAUUGUCCGGG cccca B	59
22	HCV.R1A-119 Amb.Rz-10/5	auggcucuc GagggaacucC CU UCAAGGACAUUGUCCGGG gggag B	60
23	HCV.R1A-120 Amb.Rz-10/5	uauggucuc GagggaacucC CU UCAAGGACAUUGUCCGGG cggga B	61
24	HCV.R1A-133 Amb.Rz-10/5	uuccgcagac GagggaacucC CU UCAAGGACAUUGUCCGGG acua B	62
25	HCV.R1A-140 Amb.Rz-10/5	ucaccgguuc GagggaacucC CU UCAAGGACAUUGUCCGGG gcaga B	63
26	HCV.R1A-188 Amb.Rz-10/5	cgggguuuac GagggaacucC CU UCAAGGACAUUGUCCGGG aagaa B	64
27	HCV.R1A-198 Amb.Rz-10/5	aggcaugag GagggaacucC CU UCAAGGACAUUGUCCGGG ggggu B	65
28	HCV.R1A-205 Amb.Rz-10/5	aaucuccag GagggaacucC CU UCAAGGACAUUGUCCGGG auuga B	66
29	HCV.R1A-217 Amb.Rz-10/5	ggggcacgc GagggaacucC CU UCAAGGACAUUGUCCGGG aaaa B	67
30	HCV.R1A-218 Amb.Rz-10/5	ggggcacgc GagggaacucC CU UCAAGGACAUUGUCCGGG aaau B	68
31	HCV.R1A-219 Amb.Rz-10/5	cgggggcacg GagggaacucC CU UCAAGGACAUUGUCCGGG ccaa B	69
32	HCV.R1A-223 Amb.Rz-10/5	cuugcggggg GagggaacucC CU UCAAGGACAUUGUCCGGG acgcc B	70
33	HCV.R1A-229 Amb.Rz-10/5	agcagucug GagggaacucC CU UCAAGGACAUUGUCCGGG gggg B	71
34	HCV.R1A-279 Amb.Rz-10/5	cccuacagg GagggaacucC CU UCAAGGACAUUGUCCGGG aguac B	72

Table 56

295	UGCUGCGAGUGCCCC	35	HCV.R1A-295	Amb.Rz-10/5	ggggcacucg Gaggaaaacuc CU UCAAGGACAUUCGCCGG aagca B	73
301	CGAGUGCCCCCGGAGG	36	HCV.R1A-301	Amb.Rz-10/5	ccuuccg99g Gaggaaaacuc CU UCAAGGACAUUCGCCGG acucg B	74
306	GCCCCGGGAGGUCUG	37	HCV.R1A-306	Amb.Rz-10/5	cgagaccucc Gaggaaaacuc CU UCAAGGACAUUCGCCGG g99gc B	75
307	CCCCGGGAGGUCUGU	38	HCV.R1A-307	Amb.Rz-10/5	acgagaccuc Gaggaaaacuc CU UCAAGGACAUUCGCCGG c999g B	76
No					GaaaggugugcaaccggagucaucuaauggcuuccCUUcaaggataUCgCCg	
Ribo					ggacggcB	
Ribo					GGAAAGGUGUGCAACCGGAGUCAUCAUAATGGCTCCCTUCAAGGACAUUCGCCGG	
					ACGGCB	

lower case = 2'-O-methyl
U, C = 2'-deoxy-2'-amino U, = 2'-deoxy-2'-amino C
G,A = ribo G, A
B = inverted deoxybasic

Table 57

Table 57. Additional Class II enzymatic nucleic acid Motifs

Class II Motif ID	Sequence	Seq ID No.	Kinetic Rate
A2	GGGAGGAGGAAGUGCCUGGUCAGUCACACCGAGACUGGCAGACGCUGAAACC GCCGCGCUCGCUCCCAGUCC	77	UNK
A12	GGGAGGAGGAAGUGCCUGGUGAUAUAUAUCGUUACUACGAGUGCAAGGUC GCCGCGCUCGCUCCCAGUCC	78	UNK
A11	GGGAGGAGGAAGUGCCUGGUGAGUUGCCCGAACUGUGACUACGAGUGAGGUC GCCGCGCUCGCUCCCAGUCC	79	UNK
B14	GGGAGGAGGAAGUGCCUGGCGAUCAGAUAGAGAUGAUGGCAGACGCAGAGACC GCCGCGCUCGCUCCCAGUCC	80	UNK
B10	GGGAGGAGGAAGUGCCUGGCGACUGAUACGAAAAGUCGCAGUUUCGAAACC GCCGCGCUCGCUCCCAGUCC	81	UNK
B21	GGGAGGAGGAAGUGCCUGGCGACUGAUACGAAAAGUCGCAGGUUUCGAAACC GCCGCGCUCGCUCCCAGUCC	82	UNK
B7	GGGAGGAGGAAGUGCCUUGGCUCAGCAUAAGUGAGCAGAUUGCGACACC GCCGCGCUCGCUCCCAGUCC	83	UNK
C8	GGGAGGAGGAAGUGCCUUGGUCAUUAGGAUGACAAACGUAUACUGAACACU GCCGCGCUCGCUCCCAGUCC	84	0.01 MIN ⁻¹

Table 58
Table 58: Human Her2 Class II Ribozyme and Target Sequences

RPI#	NT Pos	Substrate	Seq ID #	Ribozyme Alias	Ribozyme Sequence	Seq ID #
18722	180	CAUGGA G CUGGCG	85	erbB2-180 Zin. R2-6 amino stabl	C ₉ G ₉ C ₉ C ₉ ag GccgaaagGCGaGuccaaGGG ₉ U uccaug B	194
18835	184	GAGCUG G CGGCCU	86	erbB2-184 Zin. R2-6 amino stabl	a ₉ G ₉ G ₉ C ₉ cg GccgaaagGCGaGuccaaGGG ₉ U cagcuc B	195
18828	276	AGCUGCG G CUCCUUG	87	erbB2-276 Zin. R2-7 amino stabl	C ₉ a ₉ G ₉ G ₉ ag GccgaaagGCGaGuccaaGGG ₉ U cgcagcu B	196
18653	314	UGCUC G CCACCU	88	erbB2-314 Zin. R2-6 amino stabl	a ₉ G ₉ G ₉ u ₉ gg GccgaaagGCGaGuccaaGGG ₉ U ggaagca B	197
18825	314	AUGCUC G CCACCU	89	erbB2-314 Zin. R2-7 amino stabl	G ₉ a ₉ G ₉ G ₉ ugg GccgaaagGCGaGuccaaGGG ₉ U ggaagcau B	198
18831	379	ACCAU G CCAGCC	90	erbB2-379 Zin. R2-6 amino stabl	G ₉ G ₉ C ₉ u ₉ gg GccgaaagGCGaGuccaaGGG ₉ U auuggu B	199
18680	433	GCUAUC G CUCACAA	91	erbB2-433 Zin. R2-7 amino stabl	u ₉ u ₉ G ₉ u ₉ gag GccgaaagGCGaGuccaaGGG ₉ U gaugagc B	200
18711	594	GGAGCU G CAGCUU	92	erbB2-594 Zin. R2-6 amino stabl	a ₉ a ₉ C ₉ u ₉ g GccgaaagGCGaGuccaaGGG ₉ U agcucc B	201
18681	594	GGGAGCU G CAGCUUC	93	erbB2-594 Zin. R2-7 amino stabl	G ₉ a ₉ a ₉ G ₉ cu ₉ g GccgaaagGCGaGuccaaGGG ₉ U agcuccc B	202
18697	597	GUUGCA G CUUGCA	94	erbB2-597 Zin. R2-6 amino stabl	u ₉ C ₉ G ₉ a ₉ ag GccgaaagGCGaGuccaaGGG ₉ U ugcaagc B	203
18665	597	AGUGCA G CUUGAA	95	erbB2-597 Zin. R2-7 amino stabl	u ₉ u ₉ C ₉ a ₉ ag GccgaaagGCGaGuccaaGGG ₉ U ugcaagcu B	204
18712	659	AGUCU G CUACCA	96	erbB2-659 Zin. R2-6 amino stabl	u ₉ G ₉ G ₉ u ₉ ag GccgaaagGCGaGuccaaGGG ₉ U agagcu B	205
18682	659	CAGUCU G CUACCAG	97	erbB2-659 Zin. R2-7 amino stabl	C ₉ u ₉ G ₉ G ₉ uag GccgaaagGCGaGuccaaGGG ₉ U agagcu ₉ B	206
18683	878	CUGACU G CUGCCA	98	erbB2-878 Zin. R2-6 amino stabl	u ₉ G ₉ G ₉ C ₉ a ₉ g GccgaaagGCGaGuccaaGGG ₉ U agucag B	207
18654	878	ACUGACU G CUGCCAU	99	erbB2-878 Zin. R2-7 amino stabl	a ₉ u ₉ G ₉ G ₉ cag GccgaaagGCGaGuccaaGGG ₉ U agucag ₉ B	208
18685	881	ACUGCU G CCAUGA	100	erbB2-881 Zin. R2-6 amino stabl	u ₉ C ₉ a ₉ u ₉ gg GccgaaagGCGaGuccaaGGG ₉ U agcagu B	209
18684	881	GACUGCU G CCAUGAG	101	erbB2-881 Zin. R2-7 amino stabl	C ₉ u ₉ C ₉ a ₉ ugg GccgaaagGCGaGuccaaGGG ₉ U agcaguc B	210
18723	888	GCCAUGA G CAGUGUG	102	erbB2-888 Zin. R2-7 amino stabl	C ₉ a ₉ C ₉ a ₉ cu ₉ g GccgaaagGCGaGuccaaGGG ₉ U ucauggc B	211
18686	929	CUGACU G CCUGGC	103	erbB2-929 Zin. R2-6 amino stabl	G ₉ a ₉ C ₉ a ₉ gg GccgaaagGCGaGuccaaGGG ₉ U agucag B	212
18648	929	UCUGACU G CCUGGCC	104	erbB2-929 Zin. R2-7 amino stabl	G ₉ G ₉ C ₉ a ₉ ggg GccgaaagGCGaGuccaaGGG ₉ U agucaga B	213
18666	934	UGCCUG G CCUGCC	105	erbB2-934 Zin. R2-6 amino stabl	G ₉ G ₉ C ₉ a ₉ gg GccgaaagGCGaGuccaaGGG ₉ U caggca B	214
18651	934	CUGCCUG G CCUGCCU	106	erbB2-934 Zin. R2-7 amino stabl	a ₉ G ₉ C ₉ a ₉ ggg GccgaaagGCGaGuccaaGGG ₉ U caggcag B	215
18655	938	UGGCCU G CCUCCA	107	erbB2-938 Zin. R2-6 amino stabl	u ₉ G ₉ G ₉ a ₉ gg GccgaaagGCGaGuccaaGGG ₉ U aggccca B	216
18649	938	CUGGCCU G CCUCCAC	108	erbB2-938 Zin. R2-7 amino stabl	G ₉ u ₉ G ₉ G ₉ agg GccgaaagGCGaGuccaaGGG ₉ U aggccag B	217
18667	969	CUGUGA G CUGCAC	109	erbB2-969 Zin. R2-6 amino stabl	G ₉ u ₉ G ₉ C ₉ a ₉ g GccgaaagGCGaGuccaaGGG ₉ U ucacag B	218
18668	969	UCUGUGA G CUGCACU	110	erbB2-969 Zin. R2-7 amino stabl	a ₉ G ₉ u ₉ G ₉ cag GccgaaagGCGaGuccaaGGG ₉ U ucacaga B	219
18656	972	UGAGCU G CACUGC	111	erbB2-972 Zin. R2-6 amino stabl	G ₉ C ₉ a ₉ G ₉ u ₉ g GccgaaagGCGaGuccaaGGG ₉ U agcuca B	220
18657	972	GUAGCU G CACUGCC	112	erbB2-972 Zin. R2-7 amino stabl	G ₉ G ₉ C ₉ a ₉ du ₉ g GccgaaagGCGaGuccaaGGG ₉ U agcucac B	221
19294	972			erbB2-972 Zin. R2-6 amino stabl	G ₉ C ₉ a ₉ G ₉ u ₉ g GccaaauugGCGaGuccaaGGG ₉ U agcuca B	222

Table 58

19295	972		erbb2-972 Zin.Rz-6	amino stabl	g ₈ c ₈ a ₈ g ₈ ug GccAAuuuGgGgaGucacaaGguG ₈ u agcuca B	223
19293	972		erbb2-972 Zin.Rz-6	amino stabl	g ₈ c ₈ a ₈ g ₈ ug GccgaaagGgGgaGucagG ₈ u agcuca B	224
19292	972		erbb2-972 Zin.Rz-6	amino stabl	g ₈ c ₈ a ₈ g ₈ ug GccgaaagGgGgaGucagG ₈ u agcuca B	225
19296	972		erbb2-972 Zin.Rz-6	amino stabl	g ₈ c ₈ a ₈ g ₈ ug GccacAAuuuGgGgaGucacaaGguG ₈ u agcuca B	226
19727	972		erbb2-972 Zin.Rz-6	amino stabl	g ₈ c ₈ a ₈ g ₈ ug gccgaaagGgGgaGucagGguG ₈ u agcuca B	227
19728	972		erbb2-972 Zin.Rz-6	amino stabl	g ₈ c ₈ a ₈ g ₈ ug gccgaaagGgGgaGucagGguG ₈ u agcuca B	228
18659	1199	GAGUGU G CUAUGG	erbb2-1199 Zin.Rz-6	amino stabl	c ₈ c ₈ a ₈ u ₈ ag GccgaaagGgGgaGucacaaGguG ₈ u acacuc B	229
18658	1199	CGAGUGU G CUAUGGU	erbb2-1199 Zin.Rz-7	amino stabl	a ₈ c ₈ a ₈ u ₈ ag GccgaaagGgGgaGucacaaGguG ₈ u acacucg B	230
18724	1205	GCUAUG G UCUGGG	erbb2-1205 Zin.Rz-6	amino stabl	c ₈ c ₈ a ₈ g ₈ a GccgaaagGgGgaGucacaaGguG ₈ u cauagc B	231
18669	1205	UGCUAUG G UCUGGGC	erbb2-1205 Zin.Rz-7	amino stabl	g ₈ c ₈ a ₈ g ₈ a GccgaaagGgGgaGucacaaGguG ₈ u cauagca B	232
18725	1211	GUCUGG G CAUGGA	erbb2-1211 Zin.Rz-6	amino stabl	u ₈ c ₈ a ₈ u ₈ GccgaaagGgGgaGucacaaGguG ₈ u ccagac B	233
18726	1292	UUUGGA G CCUGGC	erbb2-1292 Zin.Rz-6	amino stabl	g ₈ c ₈ a ₈ g ₈ g GccgaaagGgGgaGucacaaGguG ₈ u ucccaa B	234
18698	1292	UUUGGA G CCUGGCA	erbb2-1292 Zin.Rz-7	amino stabl	u ₈ g ₈ c ₈ a ₈ g GccgaaagGgGgaGucacaaGguG ₈ u ucccaa B	235
18727	1313	CCGAGGA G CUUGAU	erbb2-1313 Zin.Rz-7	amino stabl	a ₈ u ₈ c ₈ a ₈ g GccgaaagGgGgaGucacaaGguG ₈ u ucuccg B	236
18699	1397	UCACAG G UUACCU	erbb2-1397 Zin.Rz-6	amino stabl	a ₈ g ₈ a ₈ u ₈ a GccgaaagGgGgaGucacaaGguG ₈ u cuguga B	237
18728	1414	AUCUA G CAUGGC	erbb2-1414 Zin.Rz-6	amino stabl	g ₈ c ₈ a ₈ u ₈ GccgaaagGgGgaGucacaaGguG ₈ u ugagau B	238
18670	1414	CAUCUA G CAUGGCC	erbb2-1414 Zin.Rz-7	amino stabl	g ₈ g ₈ c ₈ a ₈ ug GccgaaagGgGgaGucacaaGguG ₈ u ugagau B	239
18671	1536	GCUGGG G CUGGCG	erbb2-1536 Zin.Rz-6	amino stabl	g ₈ c ₈ g ₈ a ₈ g GccgaaagGgGgaGucacaaGguG ₈ u cccagc B	240
18687	1541	GGCUGC G CUCACU	erbb2-1541 Zin.Rz-6	amino stabl	a ₈ g ₈ u ₈ g ₈ ag GccgaaagGgGgaGucacaaGguG ₈ u gcagcc B	241
18829	1562	CUGGGCA G UGGACUG	erbb2-1562 Zin.Rz-7	amino stabl	c ₈ a ₈ g ₈ u ₈ GccgaaagGgGgaGucacaaGguG ₈ u ugcccag B	242
18830	1626	GGGACCA G CUCUUC	erbb2-1626 Zin.Rz-7	amino stabl	g ₈ a ₈ a ₈ g ₈ ag GccgaaagGgGgaGucacaaGguG ₈ u ugguccc B	243
18700	1755	CACCCA G UGUGUC	erbb2-1755 Zin.Rz-6	amino stabl	g ₈ a ₈ c ₈ a ₈ ca GccgaaagGgGgaGucacaaGguG ₈ u ugguug B	244
18672	1755	CCACCA G UGUGUCA	erbb2-1755 Zin.Rz-7	amino stabl	u ₈ g ₈ a ₈ c ₈ a GccgaaagGgGgaGucacaaGguG ₈ u ugguug B	245
18688	1757	CCAGU G UGUCAA	erbb2-1757 Zin.Rz-6	amino stabl	u ₈ u ₈ g ₈ a ₈ ca GccgaaagGgGgaGucacaaGguG ₈ u acuggg B	246
18660	1757	ACCCAGU G UGUCAAC	erbb2-1757 Zin.Rz-7	amino stabl	g ₈ u ₈ g ₈ a ₈ ca GccgaaagGgGgaGucacaaGguG ₈ u acuggg B	247
18689	1759	CAGUGU G UCAACU	erbb2-1759 Zin.Rz-6	amino stabl	a ₈ g ₈ u ₈ g ₈ a GccgaaagGgGgaGucacaaGguG ₈ u acacug B	248
18690	1759	CCAGUGU G UCAACUG	erbb2-1759 Zin.Rz-7	amino stabl	c ₈ a ₈ g ₈ u ₈ g ₈ a GccgaaagGgGgaGucacaaGguG ₈ u acacug B	249
18701	1784	UUUGGG G CCAGGA	erbb2-1784 Zin.Rz-6	amino stabl	u ₈ c ₈ a ₈ g ₈ g GccgaaagGgGgaGucacaaGguG ₈ u cccgaa B	250
18673	1784	CUUUGGG G CCAGGAG	erbb2-1784 Zin.Rz-7	amino stabl	c ₈ u ₈ c ₈ u ₈ g GccgaaagGgGgaGucacaaGguG ₈ u cccgaa B	251
18691	2063	UCAACU G CACCCA	erbb2-2063 Zin.Rz-6	amino stabl	u ₈ g ₈ g ₈ u ₈ g GccgaaagGgGgaGucacaaGguG ₈ u agnuga B	252
18661	2063	AUCAACU G CACCCAC	erbb2-2063 Zin.Rz-7	amino stabl	g ₈ u ₈ g ₈ g ₈ ug GccgaaagGgGgaGucacaaGguG ₈ u agnuga B	253

Table 58

18692	2075	ACUCCU G UGUCCA	138	erBB2-2075 Zin.R2-6	amino stabl	u ₉ c ₈ c ₈ ^a ₉ a GccgaagGCGaGucacagG ₉ u agga <u>g</u> u B	254
18729	2116	CAGAGA G CCAGCC	139	erBB2-2116 Zin.R2-6	amino stabl	g ₉ g ₉ c ₉ u ₉ ^{g9} GccgaagGCGaGucacagG ₉ u ucuc <u>u</u> g B	255
18832	2247	GACUGCU G CAGAAA	140	erBB2-2247 Zin.R2-7	amino stabl	u ₉ u ₉ u ₉ c ₉ ^{u9} GccgaagGCGaGucacagG ₉ u agcag <u>u</u> c B	256
18833	2271	UGGAGCC G CUACAC	141	erBB2-2271 Zin.R2-7	amino stabl	g ₉ u ₉ g ₉ u ₉ ^{c9} GccgaagGCGaGucacagG ₉ u ggcuc <u>u</u> c B	257
18702	2341	AGGAAG G UGAAGG	142	erBB2-2341 Zin.R2-6	amino stabl	c ₉ c ₉ u ₉ ^{u9} a GccgaagGCGaGucacagG ₉ u cuu <u>u</u> c B	258
18730	2347	GUGAAG G UGCUUG	143	erBB2-2347 Zin.R2-6	amino stabl	c ₉ ^a ₉ a ₉ g ₉ ^a a GccgaagGCGaGucacagG ₉ u cuue <u>a</u> c B	259
18674	2347	GGUGAAG G UGCUUGG	144	erBB2-2347 Zin.R2-7	amino stabl	c ₉ ^c ₉ a ₉ g ₉ ^a g ₉ a GccgaagGCGaGucacagG ₉ u cuue <u>a</u> c <u>c</u> B	260
18713	2349	GAAGGU G CUUGGA	145	erBB2-2349 Zin.R2-6	amino stabl	u ₉ c ₉ ^e ₉ a ₉ g ₉ GccgaagGCGaGucacagG ₉ u accu <u>u</u> c B	261
18693	2349	UGAAGGU G CUUGGAU	146	erBB2-2349 Zin.R2-7	amino stabl	a ₉ u ₉ c ₉ ^a ₉ a ₉ g ₉ GccgaagGCGaGucacagG ₉ u accu <u>u</u> c <u>a</u> B	262
18731	2384	UACAAGG G CAUCUGG	147	erBB2-2384 Zin.R2-7	amino stabl	c ₉ ^c ₉ a ₉ g ₉ ^a u ₉ GccgaagGCGaGucacagG ₉ u cu <u>u</u> g <u>u</u> a B	263
18714	2410	GGAGAAU G UGAAAUU	148	erBB2-2410 Zin.R2-7	amino stabl	a ₉ u ₉ u ₉ ^u ₉ u ₉ c ₉ a GccgaagGCGaGucacagG ₉ u auu <u>u</u> c <u>u</u> c B	264
18732	2497	GUGAUG G CUGGUG	149	erBB2-2497 Zin.R2-6	amino stabl	c ₉ ^a ₉ c ₉ ^c ₉ a ₉ g ₉ GccgaagGCGaGucacagG ₉ u cau <u>e</u> c B	265
18703	2501	UGGUG G UGUUGG	150	erBB2-2501 Zin.R2-6	amino stabl	c ₉ ^c ₉ c ₉ ^a ₉ a GccgaagGCGaGucacagG ₉ u e <u>a</u> g <u>c</u> c <u>a</u> B	266
18715	2540	GCAUCU G CTUGAC	151	erBB2-2540 Zin.R2-6	amino stabl	g ₉ u ₉ c ₉ ^a ₉ g ₉ GccgaagGCGaGucacagG ₉ u a <u>g</u> au <u>g</u> c B	267
18733	2563	CAGCUG G UGACAC	152	erBB2-2563 Zin.R2-6	amino stabl	g ₉ ^u ₉ g ₉ ^u ₉ a GccgaagGCGaGucacagG ₉ u c <u>a</u> g <u>u</u> g B	268
18734	2571	GACACA G CUUAUG	153	erBB2-2571 Zin.R2-6	amino stabl	c ₉ ^a ₉ u ₉ ^a ₉ a ₉ g ₉ GccgaagGCGaGucacagG ₉ u u <u>g</u> u <u>g</u> u <u>c</u> B	269
18675	2571	UGACACA G CUUAUGC	154	erBB2-2571 Zin.R2-7	amino stabl	g ₉ c ₉ ^a ₉ u ₉ ^a ₉ a ₉ g ₉ GccgaagGCGaGucacagG ₉ u u <u>g</u> u <u>g</u> u <u>c</u> a B	270
18716	2662	CAGAUU G CCAAGG	155	erBB2-2662 Zin.R2-6	amino stabl	c ₉ ^c ₉ u ₉ ^u ₉ g ₉ GccgaagGCGaGucacagG ₉ u a <u>a</u> cu <u>g</u> B	271
18704	2675	GGAUGA G CUACCU	156	erBB2-2675 Zin.R2-6	amino stabl	a ₉ g ₉ g ₉ ^u ₉ a ₉ g ₉ GccgaagGCGaGucacagG ₉ u u <u>u</u> cu <u>u</u> c B	272
18676	2675	GGGAUGA G CUACCUG	157	erBB2-2675 Zin.R2-7	amino stabl	c ₉ ^a ₉ g ₉ ^g ₉ a ₉ g ₉ GccgaagGCGaGucacagG ₉ u u <u>u</u> cu <u>u</u> c <u>c B</u>	273
18735	2738	GUCAAGA G UCCCAAC	158	erBB2-2738 Zin.R2-7	amino stabl	g ₉ u ₉ u ₉ ^g ₉ g ₉ a GccgaagGCGaGucacagG ₉ u u <u>u</u> cu <u>g</u> a <u>c</u> B	274
18705	2773	GGGUG G CUGGCG	159	erBB2-2773 Zin.R2-6	amino stabl	g ₉ c ₉ ^c ₉ g ₉ ^a ₉ g ₉ GccgaagGCGaGucacagG ₉ u e <u>a</u> g <u>c</u> c <u>c</u> c B	275
18836	2778	UGGUGG G CUCCUGG	160	erBB2-2778 Zin.R2-7	amino stabl	c ₉ ^c ₉ a ₉ g ₉ ^c ₉ a ₉ g ₉ GccgaagGCGaGucacagG ₉ u c <u>a</u> g <u>a</u> g <u>c</u> c <u>a</u> B	276
18694	2781	UGGGCU G CUGGAC	161	erBB2-2781 Zin.R2-6	amino stabl	g ₉ ^u ₉ c ₉ ^c ₉ a ₉ g ₉ GccgaagGCGaGucacagG ₉ u a <u>g</u> c <u>g</u> c <u>g</u> a B	277
18662	2781	CUCCGCU G CUGGACA	162	erBB2-2781 Zin.R2-7	amino stabl	u ₉ g ₉ ^u ₉ c ₉ ^a ₉ a ₉ g ₉ GccgaagGCGaGucacagG ₉ u a <u>g</u>	278
18737	2802	GACAGA G UAACAU	163	erBB2-2802 Zin.R2-6	amino stabl	a ₉ u ₉ g ₉ ^u ₉ a GccgaagGCGaGucacagG ₉ u uc <u>u</u> g <u>u</u> c B	279
18736	2802	AGACAGA G UAACAU	164	erBB2-2802 Zin.R2-7	amino stabl	c ₉ ^a ₉ u ₉ ^g ₉ a GccgaagGCGaGucacagG ₉ u uc <u>u</u> g <u>u</u> c <u>u</u> B	280
18717	2809	GUACCAU G CAGAUGG	165	erBB2-2809 Zin.R2-7	amino stabl	c ₉ ^c ₉ a ₉ u ₉ ^u ₉ c <u>u</u> g GccgaagGCGaGucacagG ₉ u au <u>g</u> g <u>u</u> a <u>c</u> B	281
18738	2819	AUGGG G CAAGGU	166	erBB2-2819 Zin.R2-6	amino stabl	a ₉ c ₉ ^c ₉ u ₉ ^u ₉ g GccgaagGCGaGucacagG ₉ u c <u>u</u> cc <u>u</u> a B	282
18706	2819	GAUGGG G CAAGGUG	167	erBB2-2819 Zin.R2-7	amino stabl	c ₉ ^a ₉ c ₉ ^c ₉ u ₉ ^u ₉ g GccgaagGCGaGucacagG ₉ u c <u>u</u> cc <u>u</u> a <u>c</u> B	283
18695	2887	GAGUAU G UGUUGAG	168	erBB2-2887 Zin.R2-7	amino stabl	c ₉ ^u ₉ c ₉ ^c ₉ a ₉ GccgaagGCGaGucacagG ₉ u au <u>a</u> cu <u>u</u> c B	284

Table 58

18663	2908	GUGACU G UGUGG	169	erBB2-2908 Zin.Rz-6 amino stabl	C ₈ C ₈ ^a C ₈ ^a ca GccgaaagGCGaGuaaGGuGū aguacac B	285
18826	2908	UGUGACU G UGUGGA	170	erBB2-2908 Zin.Rz-7 amino stabl	u ₈ C ₈ ^a C ₈ ^a ca GccgaaagGCGaGuaaGGuGū aguacac B	286
18664	2910	GACUGU G UGCGAG	171	erBB2-2910 Zin.Rz-6 amino stabl	C ₈ u ₈ C ₈ ^a ca GccgaaagGCGaGuaaGGuGū acaguc B	287
18650	2910	UGACUGU G UGCGAGC	172	erBB2-2910 Zin.Rz-7 amino stabl	G ₈ C ₈ u ₈ C ₈ ^a ca GccgaaagGCGaGuaaGGuGū acaguc B	288
18677	2916	GUGGGA G CUGAUG	173	erBB2-2916 Zin.Rz-6 amino stabl	C ₈ u ₈ u ₈ C ₈ ^a ag GccgaaagGCGaGuaaGGuGū ucccac B	289
18652	2916	UGUGGGA G CUGAUGA	174	erBB2-2916 Zin.Rz-7 amino stabl	u ₈ C ₈ ^a u ₈ C ₈ ^a ag GccgaaagGCGaGuaaGGuGū ucccac B	290
18707	2932	UUUGG G CCAAAC	175	erBB2-2932 Zin.Rz-6 amino stabl	G ₈ u ₈ u ₈ u ₈ gg GccgaaagGCGaGuaaGGuGū cccaaa B	291
18678	2932	UUUUGG G CCAAAACC	176	erBB2-2932 Zin.Rz-7 amino stabl	G ₈ u ₈ u ₈ u ₈ gg GccgaaagGCGaGuaaGGuGū cccaaa B	292
18719	3025	AUUGAU G UCUAACA	177	erBB2-3025 Zin.Rz-6 amino stabl	u ₈ u ₈ u ₈ u ₈ ga GccgaaagGCGaGuaaGGuGū aucaau B	293
18718	3025	CAUUGAU G UCUAUCAU	178	erBB2-3025 Zin.Rz-7 amino stabl	a ₈ u ₈ u ₈ u ₈ ga GccgaaagGCGaGuaaGGuGū aucaau B	294
18720	3047	UCAAAU G UUGGAU	179	erBB2-3047 Zin.Rz-6 amino stabl	a ₈ u ₈ C ₈ ^a aa GccgaaagGCGaGuaaGGuGū auuuga B	295
18696	3047	GUCAAAU G UUGGAUG	180	erBB2-3047 Zin.Rz-7 amino stabl	C ₈ u ₈ u ₈ C ₈ ^a aa GccgaaagGCGaGuaaGGuGū auuuga B	296
18739	3087	CGGGA G UUGGUG	181	erBB2-3087 Zin.Rz-6 amino stabl	C ₈ u ₈ C ₈ ^a aa GccgaaagGCGaGuaaGGuGū ucccgg B	297
18708	3087	UCCGGA G UUGGUGU	182	erBB2-3087 Zin.Rz-7 amino stabl	a ₈ C ₈ ^a C ₈ ^a aa GccgaaagGCGaGuaaGGuGū ucccgg B	298
18740	3415	GAAGG G CUGGCU	183	erBB2-3415 Zin.Rz-6 amino stabl	a ₈ u ₈ C ₈ ^a ag GccgaaagGCGaGuaaGGuGū cccuuc B	299
18741	3419	GGGUG G CUCGGA	184	erBB2-3419 Zin.Rz-6 amino stabl	u ₈ C ₈ ^a u ₈ u ₈ gg GccgaaagGCGaGuaaGGuGū cagccc B	300
18837	3419	GGGGUG G CUCGGAU	185	erBB2-3419 Zin.Rz-7 amino stabl	a ₈ u ₈ C ₈ ^a u ₈ gg GccgaaagGCGaGuaaGGuGū cagccc B	301
18709	3437	UUGAUG G UGACCU	186	erBB2-3437 Zin.Rz-6 amino stabl	a ₈ u ₈ u ₈ u ₈ ca GccgaaagGCGaGuaaGGuGū caucac B	302
18679	3437	UUUGAUG G UGACCU	187	erBB2-3437 Zin.Rz-7 amino stabl	C ₈ u ₈ u ₈ u ₈ ca GccgaaagGCGaGuaaGGuGū caucac B	303
18823	3504	UCUAACA G CCGUAC	188	erBB2-3504 Zin.Rz-6 amino stabl	G ₈ u ₈ C ₈ ^a C ₈ ^a g GccgaaagGCGaGuaaGGuGū uguaga B	304
18710	3504	CUCUAACA G CCGUAACA	189	erBB2-3504 Zin.Rz-7 amino stabl	u ₈ u ₈ u ₈ C ₈ ^a C ₈ ^a g GccgaaagGCGaGuaaGGuGū uguaga B	305
18721	3724	CAAGAC G UUUUUGC	190	erBB2-3724 Zin.Rz-7 amino stabl	G ₈ C ₈ ^a u ₈ u ₈ aa GccgaaagGCGaGuaaGGuGū gucuuug B	306
18834	3808	CCUCCU G CCUACA	191	erBB2-3808 Zin.Rz-6 amino stabl	u ₈ u ₈ u ₈ u ₈ gg GccgaaagGCGaGuaaGGuGū agggagg B	307
18827	3808	UCCUCCU G CCUACG	192	erBB2-3808 Zin.Rz-7 amino stabl	C ₈ u ₈ u ₈ u ₈ gg GccgaaagGCGaGuaaGGuGū agggagg B	308
18824	3996	GGGAG G CCUGAC	193	erBB2-3996 Zin.Rz-6 amino stabl	G ₈ u ₈ C ₈ ^a u ₈ gg GccgaaagGCGaGuaaGGuGū cuuucc B	309

UPPER CASE = RIBO

Lower case = 2'-O-methyl

C = 2'-deoxy-2'-amino Cytidine

s = phosphorothioate

B = inverted deoxyabasic

Table 59

Table 59: Human HER2 Class II (zinzyme) Ribozyme and Target Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
46	GGGCAGCC G CGGCCCC	310	GGGGCGCG GCCGAAAGGCGAGUCAAGGUCU GGCUGCCC	895
48	GCAGCCGC G CGCCCCU	311	AAGGGGCG GCCGAAAGGCGAGUCAAGGUCU GCGGCUGC	896
50	AGCCGCGC G CCCCUCC	312	GGAAGGGG GCCGAAAGGCGAGUCAAGGUCU GCGCGGCU	897
75	CCUUUACU G CGCCGCGC	313	GCGCGGCG GCCGAAAGGCGAGUCAAGGUCU AGUAAAGG	898
77	UUUACUGC G CCGCGCGC	314	GCGCGCGG GCCGAAAGGCGAGUCAAGGUCU GCAGUAAA	899
80	ACUGCGCC G CGGCCCCG	315	CGGGCGCG GCCGAAAGGCGAGUCAAGGUCU GGCAGUCU	900
82	UGCGCCGC G CGCCCGC	316	GCCGGGCG GCCGAAAGGCGAGUCAAGGUCU GCGGCGCA	901
84	CGCCGCGC G CCCGGCCC	317	GGGCCGGG GCCGAAAGGCGAGUCAAGGUCU GCGCGGCG	902
102	CACCCUUC G CAGACCC	318	GGGUGCUG GCCGAAAGGCGAGUCAAGGUCU GAGGGUG	903
112	AGCACCCC G CGCCCGC	319	GCGGGGCG GCCGAAAGGCGAGUCAAGGUCU GGGGUGCU	904
114	CACCCGCG G CCCCAGC	320	GCGCGGGG GCCGAAAGGCGAGUCAAGGUCU GCGGGGUG	905
119	CGGCCCCG G CGCCUCC	321	GGAGGGCG GCCGAAAGGCGAGUCAAGGUCU GGGGCGCG	906
121	CGCCCGCG G CCUCCCA	322	UGGGAGGG GCCGAAAGGCGAGUCAAGGUCU GCGGGGCG	907
163	CCGAGCC G CAGUGAGC	323	GCUCACUG GCCGAAAGGCGAGUCAAGGUCU GGCUCGCG	908
194	GGCCUUGU G CCGUGGG	324	CCCAGCGG GCCGAAAGGCGAGUCAAGGUCU ACAAGGCC	909
197	CUUGUGCC G CUGGGGGC	325	GCCCCCAG GCCGAAAGGCGAGUCAAGGUCU GGCACAAG	910
214	UCCUCCUC G CCUCUUG	326	CAAGAGGG GCCGAAAGGCGAGUCAAGGUCU GAGGAGGA	911
222	GCCUCUUC G CCCCCCG	327	CCGGGGGG GCCGAAAGGCGAGUCAAGGUCU AAGAGGGG	912
235	CGGAGCC G CGAGACC	328	GUUGCUG GCCGAAAGGCGAGUCAAGGUCU GGCUCGCG	913
251	CCAAGUGU G CACCGGCA	329	UGCCGUGG GCCGAAAGGCGAGUCAAGGUCU ACACUUGG	914
273	AUGAAGCU G CGGCUCCC	330	GGGAGCCG GCCGAAAGGCGAGUCAAGGUCU AGCUUCAU	915
283	GGCUCUCC G CCAGUCCC	331	GGGACUGG GCCGAAAGGCGAGUCAAGGUCU AGGGAGCC	916
309	CUGGACAU G CUCCGCCA	332	UGGCGGAG GCCGAAAGGCGAGUCAAGGUCU AUGUCCAG	917
314	CAUGCUCC G CCACCUUC	333	AGAGGUGG GCCGAAAGGCGAGUCAAGGUCU GGAGCAUG	918
332	CCAGGGCU G CCAGGUGG	334	CCACCUUG GCCGAAAGGCGAGUCAAGGUCU AGCCCUUG	919
342	CAGGUGGU G CAGGAAA	335	UUUCCUG GCCGAAAGGCGAGUCAAGGUCU ACCACUG	920
369	ACCUACCU G CCCACCA	336	UUGGUGGG GCCGAAAGGCGAGUCAAGGUCU AGGUAGGU	921
379	CCACCAAU G CCAGCCUG	337	CAGGCUUG GCCGAAAGGCGAGUCAAGGUCU AUUGGUGG	922
396	UCCUCCU G CAGGAUUA	338	AUAUCCUG GCCGAAAGGCGAGUCAAGGUCU AGGAAGGA	923
414	CAGGAGGU G CAGGGCUA	339	UAGCCUG GCCGAAAGGCGAGUCAAGGUCU ACCUCCUG	924
426	GGCUACGU G CUCAUCGC	340	GCGAUGAG GCCGAAAGGCGAGUCAAGGUCU ACGUAGCC	925
433	UGCUCauc G CUCACAAC	341	GUUGUGAG GCCGAAAGGCGAGUCAAGGUCU GAUGAGCA	926
462	GUCCACU G CAGAGGCU	342	AGCCUCUG GCCGAAAGGCGAGUCAAGGUCU AGUGGGAC	927
471	CAGAGGCU G CGGAUUGU	343	ACAAUCCG GCCGAAAGGCGAGUCAAGGUCU AGCCUCUG	928
480	CGGAUUGU G CGAGGCAC	344	GUGCCUCG GCCGAAAGGCGAGUCAAGGUCU ACAAUCCG	929
511	ACAACUUA G CCCUGGCC	345	GGCCAGGG GCCGAAAGGCGAGUCAAGGUCU AUAGUUGU	930
522	CUGGCCGU G CUAGACAA	346	UUGUCUAG GCCGAAAGGCGAGUCAAGGUCU ACGGCCAG	931
540	GGAGACCC G CUGAACAA	347	UUGUUCAG GCCGAAAGGCGAGUCAAGGUCU GGGUCUCC	932
585	GGAGGCCU G CGGGAGCU	348	AGCUCCCG GCCGAAAGGCGAGUCAAGGUCU AGGCCUCC	933
594	CGGGAGCU G CAGCUUCG	349	CGAAGCUG GCCGAAAGGCGAGUCAAGGUCU AGCUCCCG	934
659	CCAGCUCU G CUACCAGG	350	CCUGGUAG GCCGAAAGGCGAGUCAAGGUCU AGAGCUGG	935
737	CACCAACC G CUUCGGG	351	CCCAGAGG GCCGAAAGGCGAGUCAAGGUCU GGUUGGUG	936
749	UCGGGCCU G CCACCCU	352	AGGGGUGG GCCGAAAGGCGAGUCAAGGUCU AGGCCCGA	937

Table 59

782	GGGCUCCC G CUGCUGGG	353	CCCAGCAG GCCGAAAGGCGAGUCAAGGUCU	GGGAGCCC	938
785	CUCCCCGU G CUGGGGAG	354	CUCCCCAG GCCGAAAGGCGAGUCAAGGUCU	AGCGGGAG	939
822	AGCCUGAC G CGCACUGU	355	ACAGUGCG GCCGAAAGGCGAGUCAAGGUCU	GUCAGGCU	940
824	CCUGACGC G CACUGUCU	356	AGACAGUG GCCGAAAGGCGAGUCAAGGUCU	GCGUCAGG	941
835	CUGUCUGU G CCGGUGGC	357	GCCACCGG GCCGAAAGGCGAGUCAAGGUCU	ACAGACAG	942
847	GUGGCUGU G CCCGUGC	358	GCAGCGGG GCCGAAAGGCGAGUCAAGGUCU	ACAGCCAC	943
851	CUGUGCCC G CUGCAAGG	359	CCUUGCAG GCCGAAAGGCGAGUCAAGGUCU	GGGCACAG	944
854	UGCCCCGU G CAAGGGGC	360	GCCCCUUG GCCGAAAGGCGAGUCAAGGUCU	AGCGGGCA	945
867	GGGCCACU G CCCACUGA	361	UCAGUGGG GCCGAAAGGCGAGUCAAGGUCU	AGUGGCC	946
878	CACUGACU G CUGCCAUG	362	CAUGGCAG GCCGAAAGGCGAGUCAAGGUCU	AGUCAGUG	947
881	UGACUGCU G CCAUGAGC	363	GCUCAUGG GCCGAAAGGCGAGUCAAGGUCU	AGCAGUCA	948
895	AGCAGUGU G CUGCCGGC	364	GCCGGCAG GCCGAAAGGCGAGUCAAGGUCU	ACACUGCU	949
898	AGUGUGCU G CCGGUGC	365	GCAGCCGG GCCGAAAGGCGAGUCAAGGUCU	AGCACACU	950
905	UGCCGGCU G CACGGGCC	366	GGCCCGUG GCCGAAAGGCGAGUCAAGGUCU	AGCCGGCA	951
929	CUCUGACU G CCUGGCTU	367	AGGCCAGG GCCGAAAGGCGAGUCAAGGUCU	AGUCAGAG	952
938	CCUGGCCU G CCUCCACU	368	AGUGGAGG GCCGAAAGGCGAGUCAAGGUCU	AGGCCAGG	953
972	UGUGAGCU G CACUGCCC	369	GGGCAGUG GCCGAAAGGCGAGUCAAGGUCU	AGCUCACA	954
977	GCUGCACU G CCCAGCCC	370	GGGCUGGG GCCGAAAGGCGAGUCAAGGUCU	AGUGCAGC	955
1020	GAGUCCAU G CCCAUCC	371	GGAUUGGG GCCGAAAGGCGAGUCAAGGUCU	AUGGACUC	956
1051	CAUUCGGC G CCAGCUGU	372	ACAGCUGG GCCGAAAGGCGAGUCAAGGUCU	GCCGAAUG	957
1066	GUGUGACU G CCUGUCCC	373	GGGACAGG GCCGAAAGGCGAGUCAAGGUCU	AGUCACAC	958
1106	GGGAUCCU G CACCCUCG	374	CGAGGGUG GCCGAAAGGCGAGUCAAGGUCU	AGGAUCCC	959
1118	CCUCGUCU G CCCCCUGC	375	GCAGGGGG GCCGAAAGGCGAGUCAAGGUCU	AGACGAGG	960
1125	UGCCCCCU G CACAACCA	376	UGGUUGUG GCCGAAAGGCGAGUCAAGGUCU	AGGGGGCA	961
1175	UGAGAAGU G CAGCAAGC	377	GCUUGCUG GCCGAAAGGCGAGUCAAGGUCU	ACUUCUCA	962
1189	AGCCUCUG G CCCGAGUG	378	CACUCGGG GCCGAAAGGCGAGUCAAGGUCU	ACAGGGCU	963
1199	CCGAGUGU G CUAUGGUC	379	GACCAUAG GCCGAAAGGCGAGUCAAGGUCU	ACACUCGG	964
1224	GAGCACUU G CGAGAGGU	380	ACCUCUCG GCCGAAAGGCGAGUCAAGGUCU	AAGUGCUC	965
1249	UUACCAGU G CCAUAUUC	381	GAUAUUGG GCCGAAAGGCGAGUCAAGGUCU	ACUGGUAA	966
1267	AGGAGUUU G CUGGCUGC	382	GCAGCCAG GCCGAAAGGCGAGUCAAGGUCU	AAACUCCU	967
1274	UGCUGGCU G CAAGAAGA	383	UCUUCUUG GCCGAAAGGCGAGUCAAGGUCU	AGCCAGCA	968
1305	GCAUUUCU G CCGGAGAG	384	CUCUCCGG GCCGAAAGGCGAGUCAAGGUCU	AGAAAUGC	969
1342	CCAACACU G CCCCUCUC	385	GAGCGGGG GCCGAAAGGCGAGUCAAGGUCU	AGUGUUGG	970
1347	ACUGCCCC G CUCCAGCC	386	GGCUGGAG GCCGAAAGGCGAGUCAAGGUCU	GGGGCAGU	971
1431	GACAGCCU G CCUGACCU	387	AGGUCAGG GCCGAAAGGCGAGUCAAGGUCU	AGGCUGUC	972
1458	CAGAACCU G CAAGUAAU	388	AUUACUUG GCCGAAAGGCGAGUCAAGGUCU	AGGUUCUG	973
1482	CGAAUUCU G CACAAUGG	389	CCAUUGUG GCCGAAAGGCGAGUCAAGGUCU	AGAAUUCG	974
1492	ACAAUGGC G CCUACUCG	390	CGAGUAGG GCCGAAAGGCGAGUCAAGGUCU	GCCAUUGU	975
1500	GCCUACUC G CUGACCCU	391	AGGGUCAG GCCGAAAGGCGAGUCAAGGUCU	GAGUAGGC	976
1509	CUGACCCU G CAAGGGCU	392	AGCCCUUG GCCGAAAGGCGAGUCAAGGUCU	AGGGUCAG	977
1539	CUGGGGCU G CGCUCACU	393	AGUGAGCG GCCGAAAGGCGAGUCAAGGUCU	AGCCCCAG	978
1541	GGGGCUGC G CUCACUGA	394	UCAGUGAG GCCGAAAGGCGAGUCAAGGUCU	GCAGCCCC	979
1598	CCACCUCU G CUUCGUGC	395	GCACGAAG GCCGAAAGGCGAGUCAAGGUCU	AGAGGUGG	980
1605	UGCUUCGU G CACACGGU	396	ACCGUGUG GCCGAAAGGCGAGUCAAGGUCU	ACGAAGCA	981
1614	CACACGGU G CCCUGGGA	397	UCCCAGGG GCCGAAAGGCGAGUCAAGGUCU	ACCGUGUG	982
1641	CGGAACCC G CACCAAGC	398	GCUUGGUG GCCGAAAGGCGAGUCAAGGUCU	GGGUUCCG	983
1653	CAAGCUCU G CUCCACAC	399	GUGUGGAG GCCGAAAGGCGAGUCAAGGUCU	AGAGCUUG	984

Table 59

1663	UCCACACU G CCAACCGG	400	CCGGUUGG GCCGAAAGGCGAGUCAAGGUCU AGUGUGGA	985
1706	CCUGGCCU G CCACCAGC	401	GCUGGUGG GCCGAAAGGCGAGUCAAGGUCU AGGCCAGG	986
1718	CCAGCUGU G CGCCCGAG	402	CUCGGGCG GCCGAAAGGCGAGUCAAGGUCU ACAGCUGG	987
1720	AGCUGUGC G CCCGAGGG	403	CCCUCGGG GCCGAAAGGCGAGUCAAGGUCU GCACAGCU	988
1733	AGGGCACU G CUGGGGUC	404	GACCCAG GCCGAAAGGCGAGUCAAGGUCU AGUGCCCU	989
1766	UGUCAACU G CAGCCAGU	405	ACUGGCUG GCCGAAAGGCGAGUCAAGGUCU AGUUGACA	990
1793	CCAGGAGU G CGUGGAGG	406	CCUCCAG GCCGAAAGGCGAGUCAAGGUCU ACUCCUGG	991
1805	GGAGGAU G CCGAGUAC	407	GUACUCG GCCGAAAGGCGAGUCAAGGUCU AUUCCUCC	992
1815	CGAGUACU G CAGGGGCU	408	AGCCCU G GCCGAAAGGCGAGUCAAGGUCU AGUACUCG	993
1843	AUGUGAAU G CCAGGCAC	409	GUGCCUG GCCGAAAGGCGAGUCAAGGUCU AUUCACAU	994
1857	CACUGUUU G CGUGCCA	410	UGGCACG GCCGAAAGGCGAGUCAAGGUCU AAACAGUG	995
1862	UUUGCCGU G CCACCCUG	411	CAGGGUG GCCGAAAGGCGAGUCAAGGUCU ACGGCAA	996
1936	UGGCCUGU G CCCACU AU	412	AUAGUGG GCCGAAAGGCGAGUCAAGGUCU ACAGGCCA	997
1961	UCCCUUCU G CGUGGCC	413	GGGCCAG GCCGAAAGGCGAGUCAAGGUCU AGAAGGGA	998
1970	CGUGGCC G CUGCCCA	414	UGGGCAG GCCGAAAGGCGAGUCAAGGUCU GGCCACG	999
1973	GGCCCGU G CCCAGCG	415	CGCUGGG GCCGAAAGGCGAGUCAAGGUCU AGCGGGC	1000
2007	UCCUACAU G CCAUCUG	416	CAGAUGG GCCGAAAGGCGAGUCAAGGUCU AUGUAGGA	1001
2038	AGGAGGG G CAUGCCAG	417	CUGGCAUG GCCGAAAGGCGAGUCAAGGUCU GCCUCCU	1002
2042	GGGCGAU G CCAGCCU	418	AAGGCU G GCCGAAAGGCGAGUCAAGGUCU AUGCGCC	1003
2051	CCAGCCU G CCCAUCA	419	UGAUGGG GCCGAAAGGCGAGUCAAGGUCU AAGGCU G	1004
2063	CAUCAACU G CACCCACU	420	AGUGGGU GCCGAAAGGCGAGUCAAGGUCU AGUUGAU	1005
2099	CAAGGGU G CCCC GCCG	421	CGCGGGG GCCGAAAGGCGAGUCAAGGUCU AGCCCU G	1006
2104	GCUGCCCC G CCAGCAG	422	CUGCUCG GCCGAAAGGCGAGUCAAGGUCU GGGCAGC	1007
2143	UCAUCUCU G CGGUGGU	423	AACCACG GCCGAAAGGCGAGUCAAGGUCU AGAGAUGA	1008
2160	GGCAUUCU G CUGGUCU	424	ACGACCAG GCCGAAAGGCGAGUCAAGGUCU AGAAUGCC	1009
2235	UACACGAU G CGGAGACU	425	AGUCUCG GCCGAAAGGCGAGUCAAGGUCU AUCGUGUA	1010
2244	CGGAGACU G CUGCAGGA	426	UCCUCAG GCCGAAAGGCGAGUCAAGGUCU AGUCUCG	1011
2247	AGACUGCU G CAGGAAAC	427	GUUUCUG GCCGAAAGGCGAGUCAAGGUCU AGCAGUCU	1012
2271	GUGGAGCC G CUGACACC	428	GGUGUCAG GCCGAAAGGCGAGUCAAGGUCU GGCUCAC	1013
2292	GGAGCGAU G CCCAACCA	429	UGGUUGG GCCGAAAGGCGAGUCAAGGUCU AUCGCUCC	1014
2304	AACCAGG G CAGAUGCG	430	CGCAUCU GCCGAAAGGCGAGUCAAGGUCU GCCUGGU	1015
2310	GCGCAGAU G CGGAUCCU	431	AGGAUCCG GCCGAAAGGCGAGUCAAGGUCU AUCUGCG	1016
2349	GUGAAGGU G CUUGGAUC	432	GAUCCAAG GCCGAAAGGCGAGUCAAGGUCU ACCUUCAC	1017
2362	GAUCUGGC G CUUUGGC	433	GCCAAAAG GCCGAAAGGCGAGUCAAGGUCU GCCAGAU	1018
2525	UGUCUCC G CCUUCUGG	434	CCAGAAG GCCGAAAGGCGAGUCAAGGUCU GGGAGACA	1019
2540	GGGCAUCU G CCUGACAU	435	AUGUCAG GCCGAAAGGCGAGUCAAGGUCU AGAUGCCC	1020
2556	UCCACGGU G CAGCUGGU	436	ACCAGCUG GCCGAAAGGCGAGUCAAGGUCU ACCUGGA	1021
2577	CAGCUAU G CCCUAUGG	437	CCAUAGG GCCGAAAGGCGAGUCAAGGUCU AUAAGCUG	1022
2588	CUAUGGCU G CCUCUAG	438	CUAAGAG GCCGAAAGGCGAGUCAAGGUCU AGCCAUG	1023
2615	GGAAAACC G CGGACGCC	439	GGCGUCC GCCGAAAGGCGAGUCAAGGUCU GGUUUCC	1024
2621	CCGCGGAC G CCUGGGCU	440	AGCCAGG GCCGAAAGGCGAGUCAAGGUCU GUCCGCG	1025
2640	CAGGACCU G CUGAACUG	441	CAGUUCAG GCCGAAAGGCGAGUCAAGGUCU AGGUCCUG	1026
2655	UGGUGUAU G CAGAUUGC	442	GCAUUCG GCCGAAAGGCGAGUCAAGGUCU AUACACCA	1027
2662	UGCAGAUU G CCAAGGGG	443	CCCCUUG GCCGAAAGGCGAGUCAAGGUCU AAUCUGCA	1028
2691	GAGGAUGU G CGGUCUGU	444	ACGAGCCG GCCGAAAGGCGAGUCAAGGUCU ACAUCCUC	1029
2716	ACUUGGCC G CUCGGAAC	445	GUUCCAG GCCGAAAGGCGAGUCAAGGUCU GGCCAAGU	1030
2727	CGGAACGU G CUGGUCAA	446	UUGACCAG GCCGAAAGGCGAGUCAAGGUCU ACGUCCG	1031

Table 59

2781	GCUCGGCU G CUGGACAU	447	AUGUCCAG GCCGAAAGGCGAGUCAAGGUCU	AGCCGAGC	1032
2809	AGUACCAU G CAGAUGGG	448	CCCAUCUG GCCGAAAGGCGAGUCAAGGUCU	AUGGUACU	1033
2826	GGCAAGGU G CCCAUCAA	449	UUGAUGGG GCCGAAAGGCGAGUCAAGGUCU	ACCUUGCC	1034
2844	UGGAUGGC G CUGGAGUC	450	GACUCCAG GCCGAAAGGCGAGUCAAGGUCU	GCCAUCCA	1035
2861	CAUUCUCC G CCGGCGGU	451	ACCGCCGG GCCGAAAGGCGAGUCAAGGUCU	GGAGAAUG	1036
2976	CCUGACCU G CUGGAAAA	452	UUUUCCAG GCCGAAAGGCGAGUCAAGGUCU	AGGUCAGG	1037
2997	GAGCGGCU G CCCAGGCC	453	GGCUGGGG GCCGAAAGGCGAGUCAAGGUCU	AGCCGCUC	1038
3014	CCCCAUCU G CACCAUUG	454	CAAUGGUG GCCGAAAGGCGAGUCAAGGUCU	AGAUGGGG	1039
3107	AUUCUCCC G CAUGGCCA	455	UGGCCAUG GCCGAAAGGCGAGUCAAGGUCU	GGGAGAAU	1040
3128	CCCCCAGC G CUUUGUGG	456	CCACAAAG GCCGAAAGGCGAGUCAAGGUCU	GCUGGGGG	1041
3191	CUUCUACC G CUCACUGC	457	GCAGUGAG GCCGAAAGGCGAGUCAAGGUCU	GGUAGAAG	1042
3198	CGCUCACU G CUGGAGGA	458	UCCUCCAG GCCGAAAGGCGAGUCAAGGUCU	AGUGAGCG	1043
3232	UGGUGGAU G CUGAGGAG	459	CUCCUCAG GCCGAAAGGCGAGUCAAGGUCU	AUCCACCA	1044
3280	CAGACCCU G CCCCGGGC	460	GCCCGGGG GCCGAAAGGCGAGUCAAGGUCU	AGGGUCUG	1045
3289	CCCGGGC G CUGGGGGC	461	GCCCCCAG GCCGAAAGGCGAGUCAAGGUCU	GCCCGGGG	1046
3317	CAGGCACC G CAGCUCAU	462	AUGAGCUG GCCGAAAGGCGAGUCAAGGUCU	GGUGCCUG	1047
3468	AAGGGGCU G CAAAGCCU	463	AGGCUUUG GCCGAAAGGCGAGUCAAGGUCU	AGCCCCU	1048
3534	GUACCCCU G CCCUCUGA	464	UCAGAGGG GCCGAAAGGCGAGUCAAGGUCU	AGGGGUAC	1049
3559	GCUACGUU G CCCCCUG	465	CAGGGGGG GCCGAAAGGCGAGUCAAGGUCU	AACGUAGC	1050
3572	CCUGACCU G CAGCCCCC	466	GGGGGCGU GCCGAAAGGCGAGUCAAGGUCU	AGGUCAGG	1051
3627	CCCCCUUC G CCCCAGAG	467	UCUCGGGG GCCGAAAGGCGAGUCAAGGUCU	GAAGGGGG	1052
3645	GGCCCUUC G CCUGCUGC	468	GCAGCAGG GCCGAAAGGCGAGUCAAGGUCU	AGAGGGCC	1053
3649	CUCUGCCU G CUGCCCGA	469	UCGGGCAG GCCGAAAGGCGAGUCAAGGUCU	AGGCAGAG	1054
3652	UGCCUGCU G CCCGACCU	470	AGGUCGGG GCCGAAAGGCGAGUCAAGGUCU	AGCAGGCA	1055
3661	CCCGACCU G CUGGUGCC	471	GGCACCAG GCCGAAAGGCGAGUCAAGGUCU	AGGUCGGG	1056
3667	CUGCUGGU G CCACUCUG	472	CAGAGUGG GCCGAAAGGCGAGUCAAGGUCU	ACCAGCAG	1057
3730	ACGUUUUU G CCUUUGGG	473	CCCAAAGG GCCGAAAGGCGAGUCAAGGUCU	AAAAACGU	1058
3742	UUGGGGGU G CCGUGGAG	474	CUCCACGG GCCGAAAGGCGAGUCAAGGUCU	ACCCCCAA	1059
3784	GAGGAGCU G CCCUCAG	475	CUGAGGGG GCCGAAAGGCGAGUCAAGGUCU	AGCUCCUC	1060
3808	CUCCUCCU G CCUUCAGC	476	GCUGAAGG GCCGAAAGGCGAGUCAAGGUCU	AGGAGGAG	1061
3933	CUGGACGU G CCAGUGUG	477	CACACUGG GCCGAAAGGCGAGUCAAGGUCU	ACGUCCAG	1062
3960	CCAAGUCC G CAGAAGCC	478	GGCUUCUG GCCGAAAGGCGAGUCAAGGUCU	GGACUUGG	1063
4007	UGACUUCU G CUGGCAUC	479	GAUGCCAG GCCGAAAGGCGAGUCAAGGUCU	AGAAGUCA	1064
4056	GGGAACCU G CCAUGCCA	480	UGGCAUGG GCCGAAAGGCGAGUCAAGGUCU	AGGUUCCC	1065
4061	CCUGCCAU G CCAGGAAC	481	GUUCCUGG GCCGAAAGGCGAGUCAAGGUCU	AUGGCAGG	1066
4094	UCCUUCU G CUUGAGUU	482	AACUCAAG GCCGAAAGGCGAGUCAAGGUCU	AGGAAGGA	1067
4179	GAGGCCCU G CCCAUGA	483	UCAUUGGG GCCGAAAGGCGAGUCAAGGUCU	AGGGCCUC	1068
4208	CAGUGGAU G CCACAGCC	484	GGCUGUGG GCCGAAAGGCGAGUCAAGGUCU	AUCCACUG	1069
4351	CUAGUACU G CCCCCAU	485	AUGGGGGG GCCGAAAGGCGAGUCAAGGUCU	AGUACUAG	1070
4406	UACAGAGU G CUUUUCUG	486	CAGAAAAG GCCGAAAGGCGAGUCAAGGUCU	ACUCUGUA	1071
192	GCGGCCUU G UGCCGUG	487	CAGCGGCA GCCGAAAGGCGAGUCAAGGUCU	AAGGCCGC	1072
249	ACCCAAGU G UGCACGG	488	CCGGUGCA GCCGAAAGGCGAGUCAAGGUCU	ACUUGGGU	1073
387	GCCAGCCU G UCCUCCU	489	AGGAAGGA GCCGAAAGGCGAGUCAAGGUCU	AGGCUGGC	1074
478	UGCGGAUU G UGCGAGGC	490	GCCUCGCA GCCGAAAGGCGAGUCAAGGUCU	AAUCCGCA	1075
559	CCACCCCU G UCACAGG	491	CCCUGUGA GCCGAAAGGCGAGUCAAGGUCU	AGGGGUGG	1076
678	ACGAUUUU G UGGAAGGA	492	UCCUCCA GCCGAAAGGCGAGUCAAGGUCU	AAAUCGU	1077
758	CCACCCCU G UUCUCCGA	493	UCGGAGAA GCCGAAAGGCGAGUCAAGGUCU	AGGGGUGG	1078

Table 59

768	UCUCCGAU G UGUAAGGG	494	CCCUUACA GCCGAAAGGCGAGUCAAGGUCU	AUCGGAGA	1079
770	UCCGAUGU G UAAGGGCU	495	AGCCCUUA GCCGAAAGGCGAGUCAAGGUCU	ACAUCGGA	1080
809	UGAGGAU G UCAGAGCC	496	GGCUCUGA GCCGAAAGGCGAGUCAAGGUCU	AAUCCUCA	1081
829	CGCGCACU G UCUGUGCC	497	GGCACAGA GCCGAAAGGCGAGUCAAGGUCU	AGUGCGCG	1082
833	CACUGUCU G UGCCGGUG	498	CACCGGCA GCCGAAAGGCGAGUCAAGGUCU	AGACAGUG	1083
845	CGGUGGCU G UGCCCCU	499	AGCGGGCA GCCGAAAGGCGAGUCAAGGUCU	AGCCACCG	1084
893	UGAGCAGU G UGUGCCG	500	CGGCAGCA GCCGAAAGGCGAGUCAAGGUCU	ACUGCUCA	1085
965	UGGCAUCU G UGAGCUGC	501	GCAGCUCA GCCGAAAGGCGAGUCAAGGUCU	AGAUGCCA	1086
1058	CGCCAGCU G UGUGACUG	502	CAGUCACA GCCGAAAGGCGAGUCAAGGUCU	AGCUGGCG	1087
1060	CCAGCUGU G UGACUGCC	503	GGCAGUCA GCCGAAAGGCGAGUCAAGGUCU	ACAGCUGG	1088
1070	GACUGCCU G UCCCUACA	504	UGUAGGGA GCCGAAAGGCGAGUCAAGGUCU	AGGCAGUC	1089
1166	ACAGCGGU G UGAGAAU	505	ACUUCUCA GCCGAAAGGCGAGUCAAGGUCU	ACCGCUGU	1090
1187	CAAGCCCU G UGCCCCGAG	506	CUCGGGCA GCCGAAAGGCGAGUCAAGGUCU	AGGGCUUG	1091
1197	GCCCGAGU G UGCUAUGG	507	CCAUAAGCA GCCGAAAGGCGAGUCAAGGUCU	ACUCGGGC	1092
1371	CUCCAAGU G UUGAGAC	508	GUCUCAA GCCGAAAGGCGAGUCAAGGUCU	ACUUGGAG	1093
1685	GGACGAGU G UGUGGCG	509	CGCCACA GCCGAAAGGCGAGUCAAGGUCU	ACUCGUCC	1094
1687	ACGAGUGU G UGGGCGAG	510	CUCGCCCA GCCGAAAGGCGAGUCAAGGUCU	ACACUGU	1095
1716	CACCAGCU G UGCGCCG	511	CGGGCGCA GCCGAAAGGCGAGUCAAGGUCU	AGCUGGUG	1096
1757	CACCCAGU G UGUCAACU	512	AGUUGACA GCCGAAAGGCGAGUCAAGGUCU	ACUGGGUG	1097
1759	CCCAGUGU G UCAACUGC	513	GCAGUUGA GCCGAAAGGCGAGUCAAGGUCU	ACACUGGG	1098
1837	GGGAGUAU G UGAAUGCC	514	GGCAUUA GCCGAAAGGCGAGUCAAGGUCU	AUACUCCC	1099
1853	CAGGCACU G UUGCCGU	515	ACGGCAA GCCGAAAGGCGAGUCAAGGUCU	AGUGCCUG	1100
1874	CCCUGAGU G UCAGCCCC	516	GGGCGUGA GCCGAAAGGCGAGUCAAGGUCU	ACUCAGGG	1101
1901	AGUGACCU G UUUUGGAC	517	GUCCA AAA GCCGAAAGGCGAGUCAAGGUCU	AGGUCACU	1102
1925	UGACCAGU G UGUGGCU	518	AGGCCACA GCCGAAAGGCGAGUCAAGGUCU	ACUGGUCA	1103
1927	ACCAGUGU G UGGCCUGU	519	ACAGGCCA GCCGAAAGGCGAGUCAAGGUCU	ACACUGGU	1104
1934	UGUGGCU G UGCCCCU	520	AGUGGGCA GCCGAAAGGCGAGUCAAGGUCU	AGGCCACA	1105
1984	CCAGCGGU G UGAAACCU	521	AGGUUUA GCCGAAAGGCGAGUCAAGGUCU	ACCGCUGG	1106
2075	CCACUCCU G UGUGGACC	522	GGUCCACA GCCGAAAGGCGAGUCAAGGUCU	AGGAGUGG	1107
2077	ACUCCUGU G UGACCCUG	523	CAGGUCCA GCCGAAAGGCGAGUCAAGGUCU	ACAGGAGU	1108
2410	GGGAGAAU G UGAAAUAU	524	AAUUUUA GCCGAAAGGCGAGUCAAGGUCU	AUUCUCCC	1109
2436	AUCAAGU G UUGAGGGA	525	UCCCUCAA GCCGAAAGGCGAGUCAAGGUCU	ACUUUGAU	1110
2503	UGGUGGU G UGGGUCC	526	GGAGCCCA GCCGAAAGGCGAGUCAAGGUCU	ACCAGCCA	1111
2518	CCCCAUUA G UCUCGGC	527	GCGGGAGA GCCGAAAGGCGAGUCAAGGUCU	AUAUGGGG	1112
2602	UAGACCAU G UCCGGGA	528	UUCGGGA GCCGAAAGGCGAGUCAAGGUCU	AUGGUCUA	1113
2651	GAACUGGU G UAGCAGA	529	UCUGCAUA GCCGAAAGGCGAGUCAAGGUCU	ACCAGUUC	1114
2689	UGGAGGAU G UGCGGCUC	530	GAGCCGCA GCCGAAAGGCGAGUCAAGGUCU	AUCCUCCA	1115
2749	CCAACCAU G UCAAAUAU	531	AAUUUUGA GCCGAAAGGCGAGUCAAGGUCU	AUGGUUGG	1116
2887	AGAGUGAU G UGUGGAGU	532	ACUCCACA GCCGAAAGGCGAGUCAAGGUCU	AUCACUCU	1117
2889	AGUGAUGU G UGAGUUA	533	UAACUCCA GCCGAAAGGCGAGUCAAGGUCU	ACAUCACU	1118
2902	GUUAUGGU G UGACUGUG	534	CACAGUCA GCCGAAAGGCGAGUCAAGGUCU	ACCAUAAC	1119
2908	GUGUGACU G UGUGGGAG	535	CUCCACA GCCGAAAGGCGAGUCAAGGUCU	AGUCACAC	1120
2910	GUGACUGU G UGGGAGCU	536	AGCUCCA GCCGAAAGGCGAGUCAAGGUCU	ACAGUCAC	1121
3025	CCAUGAU G UCACAUG	537	CAUGUAGA GCCGAAAGGCGAGUCAAGGUCU	AUCAUUGG	1122
3047	GGUCAAU G UUGGAUGA	538	UCAUCAA GCCGAAAGGCGAGUCAAGGUCU	AUUUGACC	1123
3068	CUCUGAAU G UCGGCCAA	539	UUGGCCGA GCCGAAAGGCGAGUCAAGGUCU	AUUCAGAG	1124
3093	GAGUUGGU G UCUGAAU	540	AAUUCAGA GCCGAAAGGCGAGUCAAGGUCU	ACCAACUC	1125

Table 59

3133	AGCGCUTU G UGGUCAUC	541	GAUGACCA GCCGAAAGGCGAGUCAAGGUCU	1126
3269	CUUCUUCU G UCCAGACC	542	GGUCUGGA GCCGAAAGGCGAGUCAAGGUCU	1127
3427	GCUCCGAU G UAUUUGAU	543	AUCAAUA GCCGAAAGGCGAGUCAAGGUCU	1128
3592	CUGAAUAU G UGAACCAG	544	CUGGUUCA GCCGAAAGGCGAGUCAAGGUCU	1129
3607	AGCCAGAU G UUCGGCCC	545	GGCCCGAA GCCGAAAGGCGAGUCAAGGUCU	1130
3939	GUGCCAGU G UGAACCAG	546	CUGGUUCA GCCGAAAGGCGAGUCAAGGUCU	1131
3974	GCCCUGAU G UGUCCUCA	547	UGAGGACA GCCGAAAGGCGAGUCAAGGUCU	1132
3976	CCUGAUGU G UCCUCAGG	548	CCUGAGGA GCCGAAAGGCGAGUCAAGGUCU	1133
4072	AGGAACCU G UCCUAAGG	549	CCUAGGA GCCGAAAGGCGAGUCAAGGUCU	1134
4162	GAGUCUTU G UGGAUUCU	550	AGAAUCCA GCCGAAAGGCGAGUCAAGGUCU	1135
4300	AAGGGAGU G UCUAAGAA	551	UUCUAGA GCCGAAAGGCGAGUCAAGGUCU	1136
4332	CAGAGACU G UCCCUGAA	552	UUCAGGGA GCCGAAAGGCGAGUCAAGGUCU	1137
4380	GCAAUGGU G UCAGUAUC	553	GAUACUGA GCCGAAAGGCGAGUCAAGGUCU	1138
4397	CAGGCUTU G UACAGAGU	554	ACUCUGUA GCCGAAAGGCGAGUCAAGGUCU	1139
4414	GCUTUUCU G UTUAGUTU	555	AAACUAAA GCCGAAAGGCGAGUCAAGGUCU	1140
4434	CUUUUUUU G UTUUGUTU	556	AAACAAA GCCGAAAGGCGAGUCAAGGUCU	1141
4439	UTUUGUTU G UTUUUUUA	557	UAAAAAA GCCGAAAGGCGAGUCAAGGUCU	1142
9	AAGGGGAG G UAACCCUG	558	CAGGGUUA GCCGAAAGGCGAGUCAAGGUCU	1143
18	UAACCCUG G CCCCUUUG	559	CAAAGGGG GCCGAAAGGCGAGUCAAGGUCU	1144
27	CCCUUUG G UCGGGGCC	560	GGCCCCGA GCCGAAAGGCGAGUCAAGGUCU	1145
33	UGGUCGGG G CCCCGGGC	561	GCCCCGGG GCCGAAAGGCGAGUCAAGGUCU	1146
40	GGCCCCGG G CAGCCGCG	562	CGCGGCGU GCCGAAAGGCGAGUCAAGGUCU	1147
43	CCCGGGCA G CCGCGCGC	563	GCGCGCGG GCCGAAAGGCGAGUCAAGGUCU	1148
65	CCCACGGG G CCCUUUAC	564	GUAAAGGG GCCGAAAGGCGAGUCAAGGUCU	1149
89	CGCGCCCC G CCCCCACC	565	GGUGGGGG GCCGAAAGGCGAGUCAAGGUCU	1150
105	CCCUCCGA G CACCCCGC	566	GCGGGGUG GCCGAAAGGCGAGUCAAGGUCU	1151
130	CCCUCCCA G CCGGGUCC	567	GGACCCGG GCCGAAAGGCGAGUCAAGGUCU	1152
135	CCAGCCGG G UCCAGCCG	568	CGGCUUGA GCCGAAAGGCGAGUCAAGGUCU	1153
140	CGGGUCCA G CCGGAGCC	569	GGCUCCGG GCCGAAAGGCGAGUCAAGGUCU	1154
146	CAGCCGGA G CCAUGGGG	570	CCCCAUGG GCCGAAAGGCGAGUCAAGGUCU	1155
154	GCAUGGG G CCGGAGCC	571	GGCUCCGG GCCGAAAGGCGAGUCAAGGUCU	1156
160	GGGCCGGA G CCGCAGUG	572	CACUGCGG GCCGAAAGGCGAGUCAAGGUCU	1157
166	GAGCCGCA G UGAGCACC	573	GGUGCUCA GCCGAAAGGCGAGUCAAGGUCU	1158
170	CGCAGUGA G CACCAUGG	574	CCAUGGUG GCCGAAAGGCGAGUCAAGGUCU	1159
180	ACCAUGGA G CUGGCGGC	575	GCCGCCAG GCCGAAAGGCGAGUCAAGGUCU	1160
184	UGGAGCUG G CGGCCUUG	576	CAAGGCCG GCCGAAAGGCGAGUCAAGGUCU	1161
187	AGCUGGCG G CCUUGUGC	577	GCACAAGG GCCGAAAGGCGAGUCAAGGUCU	1162
204	CGCUGGGG G CUCCUCCU	578	AGGAGGAG GCCGAAAGGCGAGUCAAGGUCU	1163
232	CCCCCGGA G CCGCGAGC	579	GCUCGCGG GCCGAAAGGCGAGUCAAGGUCU	1164
239	AGCCGCGA G CACCCAAG	580	CUUGGGUG GCCGAAAGGCGAGUCAAGGUCU	1165
247	GCACCCAA G UGUGCACC	581	GGUGCACA GCCGAAAGGCGAGUCAAGGUCU	1166
257	GUGCACCG G CACAGACA	582	UGUCUGUG GCCGAAAGGCGAGUCAAGGUCU	1167
270	GACAUGAA G CUGCGGCU	583	AGCCGCAG GCCGAAAGGCGAGUCAAGGUCU	1168
276	AAGCUGCG G CUCCUGC	584	GCAGGGAG GCCGAAAGGCGAGUCAAGGUCU	1169
287	CCUGCCA G UCCCGAGA	585	UCUCGGGA GCCGAAAGGCGAGUCAAGGUCU	1170
329	CUACCAGG G CUGCCAGG	586	CCUGGCAG GCCGAAAGGCGAGUCAAGGUCU	1171
337	GCUGCCAG G UGGUGCAG	587	CUGCACCA GCCGAAAGGCGAGUCAAGGUCU	1172

Table 59

340	GCCAGGUG G UGCAGGGA	588	UCCCUGCA GCCGAAAGGCGAGUCAAGGUCU	CACCUGGC	1173
383	CAAUGCCA G CCUGUCCU	589	AGGACAGG GCCGAAAGGCGAGUCAAGGUCU	UGGCAUUG	1174
412	UCCAGGAG G UGCAGGGC	590	GCCCUGCA GCCGAAAGGCGAGUCAAGGUCU	CUCCUGGA	1175
419	GGUGCAGG G CUACGUGC	591	GCACGUAG GCCGAAAGGCGAGUCAAGGUCU	CCUGCACC	1176
424	AGGGCUAC G UGCUCAUC	592	GAUGAGCA GCCGAAAGGCGAGUCAAGGUCU	GUAGCCCU	1177
445	ACAACCAA G UGAGGCAG	593	CUGCCUCA GCCGAAAGGCGAGUCAAGGUCU	UUGGUUGU	1178
450	CAAGUGAG G CAGGUCCC	594	GGGACCUG GCCGAAAGGCGAGUCAAGGUCU	CUCACUUG	1179
454	UGAGGCAG G UCCCACUG	595	CAGUGGGA GCCGAAAGGCGAGUCAAGGUCU	CUGCCUCA	1180
468	CUGCAGAG G CUGCGGAU	596	AUCCGCAG GCCGAAAGGCGAGUCAAGGUCU	CUCUGCAG	1181
485	UGUGCGAG G CACCCAGC	597	GCUGGGUG GCCGAAAGGCGAGUCAAGGUCU	CUCGCACA	1182
492	GGCACCCA G CUCUUUGA	598	UCAAAGAG GCCGAAAGGCGAGUCAAGGUCU	UGGGUGCC	1183
517	AUGCCUG G CCGUGCUA	599	UAGCACGG GCCGAAAGGCGAGUCAAGGUCU	CAGGGCAU	1184
520	CCUUGGCC G UGCUAGAC	600	GUCUAGCA GCCGAAAGGCGAGUCAAGGUCU	GGCCAGGG	1185
568	UCACAGGG G CCUCCCCA	601	UGGGGAGG GCCGAAAGGCGAGUCAAGGUCU	CCCUGUGA	1186
581	CCCAGGAG G CCUGCGGG	602	CCCGCAGG GCCGAAAGGCGAGUCAAGGUCU	CUCCUGGG	1187
591	CUGCGGGA G CUGCAGCU	603	AGCUGCAG GCCGAAAGGCGAGUCAAGGUCU	UCCCGCAG	1188
597	GAGCUGCA G CUUCGAAG	604	CUUCGAAG GCCGAAAGGCGAGUCAAGGUCU	UGCAGCUC	1189
605	GCUUCGAA G CCUCACAG	605	CUGUGAGG GCCGAAAGGCGAGUCAAGGUCU	UUCGAAAGC	1190
631	AAGGAGGG G UCUUGAUC	606	GAUCAAGA GCCGAAAGGCGAGUCAAGGUCU	CCCUCCUU	1191
642	UUGAUCCA G CGGAACCC	607	GGGUUCCG GCCGAAAGGCGAGUCAAGGUCU	UGGAUCAA	1192
654	AACCCCCA G CUCUGCUA	608	UAGCAGAG GCCGAAAGGCGAGUCAAGGUCU	UGGGGGUU	1193
708	AACAACCA G CUGGCUCU	609	AGAGCCAG GCCGAAAGGCGAGUCAAGGUCU	UGGUUGUU	1194
712	ACCAGCUG G CUCUCACA	610	UGUGAGAG GCCGAAAGGCGAGUCAAGGUCU	CAGCUGGU	1195
745	GCUCUCGG G CCUGCCAC	611	GUGGCAGG GCCGAAAGGCGAGUCAAGGUCU	CCGAGAGC	1196
776	GUGUAAGG G CUCCCGCU	612	AGCGGGAG GCCGAAAGGCGAGUCAAGGUCU	CCUUAACAC	1197
797	GGGAGAGA G UUCUGAGG	613	CCUCAGAA GCCGAAAGGCGAGUCAAGGUCU	UTCUCUCC	1198
815	UUGUCAGA G CCUGACGC	614	GCGUCAGG GCCGAAAGGCGAGUCAAGGUCU	UCUGACAA	1199
839	CUGUGCCG G UGGCUGUG	615	CACAGCCA GCCGAAAGGCGAGUCAAGGUCU	CGGCACAG	1200
842	UGCCGGUG G CUGUGCCC	616	GGGCACAG GCCGAAAGGCGAGUCAAGGUCU	CACCGGCA	1201
861	UGCAAGGG G CCACUGCC	617	GGCAGUGG GCCGAAAGGCGAGUCAAGGUCU	CCCUUGCA	1202
888	UGCCAUGA G CAGUGUGC	618	GCACACUG GCCGAAAGGCGAGUCAAGGUCU	UCAUGGCA	1203
891	CAUGAGCA G UGUGCUGC	619	GCAGCACA GCCGAAAGGCGAGUCAAGGUCU	UGCUC AUG	1204
902	UGCUGCCG G CUGCACGG	620	CCGUGCAG GCCGAAAGGCGAGUCAAGGUCU	CGGCAGCA	1205
911	CUGCACGG G CCCCAGC	621	GCUUGGGG GCCGAAAGGCGAGUCAAGGUCU	CCGUGCAG	1206
918	GGCCCCAA G CACUCUGA	622	UCAGAGUG GCCGAAAGGCGAGUCAAGGUCU	UUGGGGCC	1207
934	ACUGCCUG G CCUGCCUC	623	GAGGCAGG GCCGAAAGGCGAGUCAAGGUCU	CAGGCAGU	1208
956	CAACCACA G UGGCAUCU	624	AGAUGCCA GCCGAAAGGCGAGUCAAGGUCU	UGUGGUUG	1209
959	CCACAGUG G CAUCUGUG	625	CACAGAUG GCCGAAAGGCGAGUCAAGGUCU	CACUGUGG	1210
969	AUCUGUGA G CUGCACUG	626	CAGUGCAG GCCGAAAGGCGAGUCAAGGUCU	UCACAGAU	1211
982	ACUGCCCA G CCCUGGUC	627	GACCAGGG GCCGAAAGGCGAGUCAAGGUCU	UGGGCAGU	1212
988	CAGCCUG G UCACCUAC	628	GUAGGUGA GCCGAAAGGCGAGUCAAGGUCU	CAGGGCUG	1213
1008	ACAGACAC G UUUGAGUC	629	GACUCAAA GCCGAAAGGCGAGUCAAGGUCU	GUGUCUGU	1214
1014	ACGUUUGA G UCCAUGCC	630	GGCAUGGA GCCGAAAGGCGAGUCAAGGUCU	UCAAACGU	1215
1034	UCCCGAGG G CCGUAUA	631	UAUACCGG GCCGAAAGGCGAGUCAAGGUCU	CCUCGGGA	1216
1038	GAGGGCCG G UAUACAUU	632	AAUGUAUA GCCGAAAGGCGAGUCAAGGUCU	CGGCCUC	1217
1049	UACAUUCG G CGCCAGCU	633	AGCUGGCG GCCGAAAGGCGAGUCAAGGUCU	CGAAUGUA	1218
1055	CGGCGCCA G CUGUGUGA	634	UCACACAG GCCGAAAGGCGAGUCAAGGUCU	UGGCGCCG	1219

Table 59

1096	CUACGGAC G UGGGAUCC	635	GAUCCCA GCCGAAAGGCGAGUCAAGGUCU GUCCGUAG	1220
1114	GCACCCUC G UCUGCCCC	636	GGGGCAGA GCCGAAAGGCGAGUCAAGGUCU GAGGGUGC	1221
1138	ACCAAGAG G UGACAGCA	637	UGCUGUCA GCCGAAAGGCGAGUCAAGGUCU CUCUUGGU	1222
1144	AGGUGACA G CAGAGGAU	638	AUCCUCUG GCCGAAAGGCGAGUCAAGGUCU UGUCACCU	1223
1161	GGAACACA G CGUGUGA	639	UCACACCG GCCGAAAGGCGAGUCAAGGUCU UGUGUCC	1224
1164	ACACAGCG G UGUGAGAA	640	UUCUCACA GCCGAAAGGCGAGUCAAGGUCU CGCUGUGU	1225
1173	UGUGAGAA G UGCAGCAA	641	UUGCUGCA GCCGAAAGGCGAGUCAAGGUCU UUCUCACA	1226
1178	GAAGUGCA G CAAGCCU	642	AGGGCUUG GCCGAAAGGCGAGUCAAGGUCU UGCACUUC	1227
1182	UGCAGCAA G CCCUGUGC	643	GCACAGGG GCCGAAAGGCGAGUCAAGGUCU UUGCUGCA	1228
1195	GUGCCCGA G UGUGCUAU	644	AUAGCACA GCCGAAAGGCGAGUCAAGGUCU UCGGGCAC	1229
1205	GUGCUAUG G UCUGGCA	645	UGCCAGA GCCGAAAGGCGAGUCAAGGUCU CAUAGCAC	1230
1211	UGGUCUGG G CAUGGAGC	646	GUCCAUUG GCCGAAAGGCGAGUCAAGGUCU CCAGACCA	1231
1218	GGCAUGGA G CACUUGCG	647	CGCAAGUG GCCGAAAGGCGAGUCAAGGUCU UCCAUGCC	1232
1231	UGCGAGAG G UGAGGGCA	648	UGCCUCA GCCGAAAGGCGAGUCAAGGUCU CUCUCGCA	1233
1237	AGGUGAGG G CAGUACC	649	GGUACUG GCCGAAAGGCGAGUCAAGGUCU CCUCACCU	1234
1240	UGAGGGCA G UUACCAGU	650	ACUGGUAA GCCGAAAGGCGAGUCAAGGUCU UGCCCUGA	1235
1247	AGUUAACA G UGCCAAUA	651	UAUUGGCA GCCGAAAGGCGAGUCAAGGUCU UGGUAAU	1236
1263	AUCCAGGA G UUGCUGG	652	CCAGCAA GCCGAAAGGCGAGUCAAGGUCU UCCUGAU	1237
1271	GUUUGCUG G CUGCAAGA	653	UCUUGCAG GCCGAAAGGCGAGUCAAGGUCU CAGCAAAC	1238
1292	CUUUGGA G CCUGGCAU	654	AUGCCAG GCCGAAAGGCGAGUCAAGGUCU UCCCAAAG	1239
1297	GGAGCCUG G CAUUCUG	655	CAGAAUG GCCGAAAGGCGAGUCAAGGUCU CAGGCUC	1240
1313	GCCGAGA G CUUGAUG	656	CAUCAAAG GCCGAAAGGCGAGUCAAGGUCU UCUCGGC	1241
1330	GGGACCA G CCUCCAA	657	GUUGGAG GCCGAAAGGCGAGUCAAGGUCU UGGGUCC	1242
1353	CCGCUCA G CCAGAGCA	658	UGCUCUG GCCGAAAGGCGAGUCAAGGUCU UGGAGCG	1243
1359	CAGCCAGA G CAGCUCA	659	UGGAGCUG GCCGAAAGGCGAGUCAAGGUCU UCUGGCUG	1244
1362	CCAGAGCA G CUCCAAGU	660	ACUUGGAG GCCGAAAGGCGAGUCAAGGUCU UGCUCUG	1245
1369	AGCUCCA G UGUUGAG	661	CUCAAACA GCCGAAAGGCGAGUCAAGGUCU UUGGAGCU	1246
1397	GAUCACAG G UUAUUAU	662	AUAGGUAA GCCGAAAGGCGAGUCAAGGUCU CUGUGAU	1247
1414	ACAUCUA G CAUGGCCG	663	CGGCCAUG GCCGAAAGGCGAGUCAAGGUCU UGAGAUGU	1248
1419	UCAGCAUG G CCGACAG	664	CUGUCCG GCCGAAAGGCGAGUCAAGGUCU CAUGCUGA	1249
1427	GCCGACA G CCUGCCUG	665	CAGGCAG GCCGAAAGGCGAGUCAAGGUCU UGUCCGC	1250
1442	UGACCUCA G CGUCUCC	666	GGAGACG GCCGAAAGGCGAGUCAAGGUCU UGAGGUCA	1251
1444	ACCUCAG G UCUCACAG	667	CUGGAAGA GCCGAAAGGCGAGUCAAGGUCU GCUGAGGU	1252
1462	ACCUGCA G UAAUCCG	668	CCGGAUA GCCGAAAGGCGAGUCAAGGUCU UUGCAGGU	1253
1490	GCACAAUG G CGCCUACU	669	AGUAGGCG GCCGAAAGGCGAGUCAAGGUCU CAUUGUG	1254
1515	CUGCAAGG G CUGGGCAU	670	AUGCCCAG GCCGAAAGGCGAGUCAAGGUCU CCUUGCAG	1255
1520	AGGGCUGG G CAUCAGCU	671	AGCUGAUG GCCGAAAGGCGAGUCAAGGUCU CCAGCCU	1256
1526	GGGCAUA G CUGGCUGG	672	CCAGCCAG GCCGAAAGGCGAGUCAAGGUCU UGAUGCC	1257
1530	AUCAGCUG G CUGGGCU	673	AGCCCCAG GCCGAAAGGCGAGUCAAGGUCU CAGCUGAU	1258
1536	UGGUGGG G CUGGCUC	674	GAGCGCAG GCCGAAAGGCGAGUCAAGGUCU CCCAGCCA	1259
1559	GGAACUGG G CAGUGGAC	675	GUCCACUG GCCGAAAGGCGAGUCAAGGUCU CCAGUCC	1260
1562	ACUGGGCA G UGGACUGG	676	CCAGUCCA GCCGAAAGGCGAGUCAAGGUCU UGCCCAGU	1261
1570	GUGGACUG G CCCUCAUC	677	GAUGAGGG GCCGAAAGGCGAGUCAAGGUCU CAGUCCAC	1262
1603	UCUGCUUC G UGCACAG	678	CGUGUGCA GCCGAAAGGCGAGUCAAGGUCU GAAGCAGA	1263
1612	UGCACAG G UGCCCUGG	679	CCAGGGCA GCCGAAAGGCGAGUCAAGGUCU CGUGUGCA	1264
1626	UGGGACCA G CUCUUCG	680	CGAAAGAG GCCGAAAGGCGAGUCAAGGUCU UGGUCCA	1265
1648	CGCACCA G CUCUGCUC	681	GAGCAGAG GCCGAAAGGCGAGUCAAGGUCU UUGGUGCG	1266

Table 59

1671	GCCAACCG G CCAGAGGA	682	UCCUCUGG GCCGAAAGGCGAGUCAAGGUCU CGGUUGGC	1267
1683	GAGGACGA G UGUGUGGG	683	CCCACACA GCCGAAAGGCGAGUCAAGGUCU UCGUCCUC	1268
1691	GUGUGUGG G CGAGGGCC	684	GGCCUCUG GCCGAAAGGCGAGUCAAGGUCU CCACACAC	1269
1697	GGGCGAGG G CCUGGCCU	685	AGGCCAGG GCCGAAAGGCGAGUCAAGGUCU CCUCGCCC	1270
1702	AGGGCCUG G CCUGCCAC	686	GUGGCAGG GCCGAAAGGCGAGUCAAGGUCU CAGGCCCU	1271
1713	UGCCACCA G CUGUGCGC	687	GCGCACAG GCCGAAAGGCGAGUCAAGGUCU UGGUGGCA	1272
1728	GCCCGAGG G CACUGCUG	688	CAGCAGUG GCCGAAAGGCGAGUCAAGGUCU CCUCGGGC	1273
1739	CUGCUGGG G UCCAGGGC	689	GCCCUGGA GCCGAAAGGCGAGUCAAGGUCU CCCAGCAG	1274
1746	GGUCCAGG G CCCACCCA	690	UGGGUGGG GCCGAAAGGCGAGUCAAGGUCU CCUGGACC	1275
1755	CCCACCCA G UGUGUCAA	691	UUGACACA GCCGAAAGGCGAGUCAAGGUCU UGGGUGGG	1276
1769	CAACUGCA G CCAGUUC	692	GGAACUGG GCCGAAAGGCGAGUCAAGGUCU UGCAGUUG	1277
1773	UGCAGCCA G UUCUUCG	693	CGAAGGAA GCCGAAAGGCGAGUCAAGGUCU UGGCUGCA	1278
1784	CCUUCGGG G CCAGGAGU	694	ACUCCUGG GCCGAAAGGCGAGUCAAGGUCU CCCGAAGG	1279
1791	GGCCAGGA G UGCGUGGA	695	UCCACGCA GCCGAAAGGCGAGUCAAGGUCU UCCUGGCC	1280
1795	AGGAGUGC G UGGAGGAA	696	UUCCUCCA GCCGAAAGGCGAGUCAAGGUCU GCACUCCU	1281
1810	AAUGCCGA G UACUGCAG	697	CUGCAGUA GCCGAAAGGCGAGUCAAGGUCU UCGGCAUU	1282
1821	CUGCAGGG G CUCCCCAG	698	CUGGGGAG GCCGAAAGGCGAGUCAAGGUCU CCCUGCAG	1283
1833	CCCAGGGA G UAUGUGAA	699	UUCACAUU GCCGAAAGGCGAGUCAAGGUCU UCCCUGGG	1284
1848	AAUGCCAG G CACUGUUU	700	AAACAGUG GCCGAAAGGCGAGUCAAGGUCU CUGGCAUU	1285
1860	UGUUUGCC G UGCCACCC	701	GGGUGGCA GCCGAAAGGCGAGUCAAGGUCU GGCAAACA	1286
1872	CACCCUGA G UGUCAGCC	702	GGCUGACA GCCGAAAGGCGAGUCAAGGUCU UCAGGGUG	1287
1878	GAGUGUCA G CCCAGAA	703	UUCUGGGG GCCGAAAGGCGAGUCAAGGUCU UGACACUC	1288
1889	CCAGAAUG G CUCAGUGA	704	UCACUGAG GCCGAAAGGCGAGUCAAGGUCU CAUUCUGG	1289
1894	AUGGCUCA G UGACCUGU	705	ACAGGUCA GCCGAAAGGCGAGUCAAGGUCU UGAGCAU	1290
1915	GACCGGAG G CUGACCAG	706	CUGGUCAG GCCGAAAGGCGAGUCAAGGUCU CUCCGGUC	1291
1923	GCUGACCA G UGUGUGGC	707	GCCACACA GCCGAAAGGCGAGUCAAGGUCU UGGUCAGC	1292
1930	AGUGUGUG G CCUGUGCC	708	GGCACAGG GCCGAAAGGCGAGUCAAGGUCU CACACACU	1293
1963	CCUUCUGC G UGGCCCGC	709	GCGGGCCA GCCGAAAGGCGAGUCAAGGUCU GCAGAAGG	1294
1966	UCUGCGUG G CCGCUGC	710	GCAGCGGG GCCGAAAGGCGAGUCAAGGUCU CACGCAGA	1295
1979	CUGCCCCA G CGGUGUGA	711	UCACACCG GCCGAAAGGCGAGUCAAGGUCU UGGGGCAG	1296
1982	CCCCAGCG G UGUGAAAC	712	GUUUCACA GCCGAAAGGCGAGUCAAGGUCU CGCUGGGG	1297
2019	AUCUGGAA G UUUCCAGA	713	UCUGGAAA GCCGAAAGGCGAGUCAAGGUCU UUCCAGAU	1298
2036	UGAGGAGG G CGCAUGCC	714	GGCAUGCG GCCGAAAGGCGAGUCAAGGUCU CCUCCUCA	1299
2046	GCAUGCCA G CCUUGCCC	715	GGGCAAGG GCCGAAAGGCGAGUCAAGGUCU UGGCAUGC	1300
2096	UGACAAGG G CUGCCCCG	716	CGGGGCAG GCCGAAAGGCGAGUCAAGGUCU CCUUGUCA	1301
2109	CCCGCCGA G CAGAGAGC	717	GCUCUCUG GCCGAAAGGCGAGUCAAGGUCU UCGGCGGG	1302
2116	AGCAGAGA G CCAGCCCU	718	AGGGCUGG GCCGAAAGGCGAGUCAAGGUCU UCUCUGCU	1303
2120	GAGAGCCA G CCCUCUGA	719	UCAGAGGG GCCGAAAGGCGAGUCAAGGUCU UGGCUCUC	1304
2130	CCUCUGAC G UCCAUCAU	720	AUGAUGGA GCCGAAAGGCGAGUCAAGGUCU GUCAGAGG	1305
2146	UCUCUGCG G UGGUUGGC	721	GCCAACCA GCCGAAAGGCGAGUCAAGGUCU CGCAGAGA	1306
2149	CUGCGGUG G UUGGCAUU	722	AAUGCCAA GCCGAAAGGCGAGUCAAGGUCU CACCGCAG	1307
2153	GGUGGUUG G CAUUCUGC	723	GCAGAAUG GCCGAAAGGCGAGUCAAGGUCU CAACCACC	1308
2164	UUCUGCUG G UCGUGGUC	724	GACCACGA GCCGAAAGGCGAGUCAAGGUCU CAGCAGAA	1309
2167	UGCUGGUC G UGGUCUUG	725	CAAGACCA GCCGAAAGGCGAGUCAAGGUCU GACCAGCA	1310
2170	UGGUCGUG G UCUGGGG	726	CCCCAAGA GCCGAAAGGCGAGUCAAGGUCU CACGACCA	1311
2179	UCUUGGGG G UGGUCUUU	727	AAAGACCA GCCGAAAGGCGAGUCAAGGUCU CCCCAGA	1312
2182	UGGGGGUG G UCUUGGG	728	CCCAAAGA GCCGAAAGGCGAGUCAAGGUCU CACCCCA	1313

Table 59

2202	CUCAUCAA G CGACGGCA	729	UGCCGUCG GCCGAAAGGCGAGUCAAGGUCU UUGAUGAG	1314
2208	AAGCGACG G CAGCAGAA	730	UUCUGCUG GCCGAAAGGCGAGUCAAGGUCU CGUCGCUU	1315
2211	CGACGGCA G CAGAAGAU	731	AUCUUCUG GCCGAAAGGCGAGUCAAGGUCU UGCCGUCG	1316
2226	AUCCGGAA G UACACGAU	732	AUCGUGUA GCCGAAAGGCGAGUCAAGGUCU UUCGGAU	1317
2259	GAAACGGA G CUGGUGGA	733	UCCACCAG GCCGAAAGGCGAGUCAAGGUCU UCCGUUUC	1318
2263	CGGAGCUG G UGGAGCCG	734	CGGCUCCA GCCGAAAGGCGAGUCAAGGUCU CAGCUCCG	1319
2268	CUGGUGGA G CCGCUGAC	735	GUCAGCGG GCCGAAAGGCGAGUCAAGGUCU UCCACCAG	1320
2282	GACACCUA G CGGAGCGA	736	UCGCUCCG GCCGAAAGGCGAGUCAAGGUCU UAGGUGUC	1321
2287	CUAGCGGA G CGAUGCCC	737	GGGCAUCG GCCGAAAGGCGAGUCAAGGUCU UCCGCUAG	1322
2302	CCAACCAG G CGCAGAUG	738	CAUCUGCG GCCGAAAGGCGAGUCAAGGUCU CUGGUUGG	1323
2331	GAGACGGA G CUGAGGAA	739	UUCUCAG GCCGAAAGGCGAGUCAAGGUCU UCCGUCUC	1324
2341	UGAGGAAG G UGAAGGUG	740	CACCUUCA GCCGAAAGGCGAGUCAAGGUCU CUUCCUCA	1325
2347	AGGUGAAG G UGCUUGGA	741	UCCAAGCA GCCGAAAGGCGAGUCAAGGUCU CUUACCU	1326
2360	UGGAUCUG G CGCUUUG	742	CAAAAGCG GCCGAAAGGCGAGUCAAGGUCU CAGAUCCA	1327
2369	CGCUUUG G CACAGUCU	743	AGACUGUG GCCGAAAGGCGAGUCAAGGUCU CAAAAGCG	1328
2374	UUGGCACA G UCUACAAG	744	CUUGUAGA GCCGAAAGGCGAGUCAAGGUCU UGUGCCAA	1329
2384	CUACAAGG G CAUCUGGA	745	UCCAGAUG GCCGAAAGGCGAGUCAAGGUCU CCUGUAG	1330
2422	AAAUUCCA G UGGCCAUC	746	GAUGGCCA GCCGAAAGGCGAGUCAAGGUCU UGGAUUU	1331
2425	UUCACAG G CCAUCAA	747	UUUGAUGG GCCGAAAGGCGAGUCAAGGUCU CACUGGAA	1332
2434	CCAUCAA G UGUUGAGG	748	CCUCAACA GCCGAAAGGCGAGUCAAGGUCU UUUGAUGG	1333
2461	CCCCCAA G CCAACAA	749	UUUGUUGG GCCGAAAGGCGAGUCAAGGUCU UUUGGGG	1334
2485	UAGACGAA G CAUACGUG	750	CACGUAUG GCCGAAAGGCGAGUCAAGGUCU UUCGUCUA	1335
2491	AAGCAUAC G UGAUGGCU	751	AGCCAUCA GCCGAAAGGCGAGUCAAGGUCU GUUUGCUU	1336
2497	ACGUGAUG G CUGGUGUG	752	CACACCAG GCCGAAAGGCGAGUCAAGGUCU CAUACGUG	1337
2501	GAUGGCUG G UGUGGCU	753	AGCCACA GCCGAAAGGCGAGUCAAGGUCU CAGCCAUC	1338
2507	UGGUGUGG G CUCCCAU	754	AUGGGGAG GCCGAAAGGCGAGUCAAGGUCU CCACACCA	1339
2534	CCUUCUGG G CAUCUGCC	755	GGCAGAUG GCCGAAAGGCGAGUCAAGGUCU CCAGAAGG	1340
2554	CAUCCACG G UGCAGCUG	756	CAGCUGCA GCCGAAAGGCGAGUCAAGGUCU CGUGGAUG	1341
2559	ACGUGUCA G CUGGUGAC	757	GUCACCAG GCCGAAAGGCGAGUCAAGGUCU UGCACCGU	1342
2563	UGCAGCUG G UGACACAG	758	CUGUGUCA GCCGAAAGGCGAGUCAAGGUCU CAGCUGCA	1343
2571	GUGACACA G CUUAUGCC	759	GGCAUAAG GCCGAAAGGCGAGUCAAGGUCU UGUGUCAC	1344
2585	GCCCUAUG G CUGCCUCU	760	AGAGGCAG GCCGAAAGGCGAGUCAAGGUCU CAUAGGGC	1345
2627	ACGCCUGG G CUCCAGG	761	CCUGGGAG GCCGAAAGGCGAGUCAAGGUCU CCAGGCGU	1346
2649	CUGAACUG G UGUUUGCA	762	UGCAUACA GCCGAAAGGCGAGUCAAGGUCU CAGUUCAG	1347
2675	GGGAUGA G CUACCUUG	763	CCAGGUAG GCCGAAAGGCGAGUCAAGGUCU UCAUCCCC	1348
2694	GAUGUGCG G CUCGUACA	764	UGUACGAG GCCGAAAGGCGAGUCAAGGUCU CGCACAU	1349
2698	UGCUGCUC G UACACAGG	765	CCUGUGUA GCCGAAAGGCGAGUCAAGGUCU GAGCCGCA	1350
2713	GGGACUUG G CCGCUCGG	766	CCGAGCGG GCCGAAAGGCGAGUCAAGGUCU CAAGUCCC	1351
2725	CUCGGAAC G UGCUGGUC	767	GACCAGCA GCCGAAAGGCGAGUCAAGGUCU GUUCCGAG	1352
2731	ACGUGCUG G UCAAGAGU	768	ACUCUUGA GCCGAAAGGCGAGUCAAGGUCU CAGCACGU	1353
2738	GUCAAGA G UCCCAACC	769	GGUUGGGA GCCGAAAGGCGAGUCAAGGUCU UCUGAGCC	1354
2769	GACUUCGG G CUGGCUCG	770	CGAGCCAG GCCGAAAGGCGAGUCAAGGUCU CCGAAGUC	1355
2773	UGGGCUG G CUCGGCUG	771	CAGCCGAG GCCGAAAGGCGAGUCAAGGUCU CAGCCCGA	1356
2778	CUGGCUCG G CUGCUGGA	772	UCCAGCAG GCCGAAAGGCGAGUCAAGGUCU CGAGCCAG	1357
2802	GAGACAGA G UACCAUGC	773	GCAUGGUA GCCGAAAGGCGAGUCAAGGUCU UCUGUCUC	1358
2819	AGAUGGGG G CAAGGUGC	774	GCACCUUG GCCGAAAGGCGAGUCAAGGUCU CCCCUCU	1359
2824	GGGCAAG G UGCCCAUC	775	GAUGGGCA GCCGAAAGGCGAGUCAAGGUCU CUUGCCCC	1360

Table 59

2835	CCCAUCAA G UGGAUGGC	776	GCCAUCCA GCCGAAAGGCGAGUCAAGGUCU UUGAUGGG	1361
2842	AGUGGAUG G CGCUGGAG	777	CUCCAGCG GCCGAAAGGCGAGUCAAGGUCU CAUCCACU	1362
2850	GCGCUGGA G UCCAUUCU	778	AGAAUGGA GCCGAAAGGCGAGUCAAGGUCU UCCAGCGC	1363
2865	CUCCGCCG G CGGUUCAC	779	GUGAACCG GCCGAAAGGCGAGUCAAGGUCU CGCGGGAG	1364
2868	CGCCGGCG G UUCACCCA	780	UGGGUGAA GCCGAAAGGCGAGUCAAGGUCU CGCCGGCG	1365
2882	CCACCAGA G UGAUGUGU	781	ACACAUCA GCCGAAAGGCGAGUCAAGGUCU UCUGGUGG	1366
2894	UGUGUGGA G UUAUGGUG	782	CACCAUAA GCCGAAAGGCGAGUCAAGGUCU UCCACACA	1367
2900	GAGUUAUG G UGUGACUG	783	CAGUCACA GCCGAAAGGCGAGUCAAGGUCU CAUAACUC	1368
2916	GUGUGGGA G CUGAUGAC	784	GUCAUCAG GCCGAAAGGCGAGUCAAGGUCU UCCCACAC	1369
2932	CUUUUGGG G CCAAACCU	785	AGGUUUGG GCCGAAAGGCGAGUCAAGGUCU CCCAAAG	1370
2956	GAUCCCA G CCCGGGAG	786	CUCCCGGG GCCGAAAGGCGAGUCAAGGUCU UGGGAUCC	1371
2991	AAGGGGGA G CGGCUGCC	787	GGCAGCCG GCCGAAAGGCGAGUCAAGGUCU UCCCCU	1372
2994	GGGAGCG G CUGCCCCA	788	UGGGGCGAG GCCGAAAGGCGAGUCAAGGUCU CGCUCCCC	1373
3003	CUGCCCCA G CCCCCAU	789	AUGGGGGG GCCGAAAGGCGAGUCAAGGUCU UGGGGCAG	1374
3040	UGAUCAUG G UCAAAUGU	790	ACAUUUGA GCCGAAAGGCGAGUCAAGGUCU CAUGAUCA	1375
3072	GAAUGUCG G CCAAGAUU	791	AAUCUUGG GCCGAAAGGCGAGUCAAGGUCU CGACAUUC	1376
3087	UUCCGGGA G UUGGUGUC	792	GACACCAA GCCGAAAGGCGAGUCAAGGUCU UCCCGGAA	1377
3091	GGGAGUUG G UGUCUGAA	793	UUCAGACA GCCGAAAGGCGAGUCAAGGUCU CAACUCCC	1378
3112	CCCGCAUG G CCAGGGAC	794	GUCCUGG GCCGAAAGGCGAGUCAAGGUCU CAUGCGGG	1379
3126	GACCCCCA G CGCUUUGU	795	ACAAAGCG GCCGAAAGGCGAGUCAAGGUCU UGGGGGUC	1380
3136	GCUUUGUG G UCAUCCAG	796	CUGGAUGA GCCGAAAGGCGAGUCAAGGUCU CACAAAGC	1381
3158	GGACUUGG G CCCAGCCA	797	UGGCUGGG GCCGAAAGGCGAGUCAAGGUCU CCAAGUCC	1382
3163	UGGGCCCA G CAGUCCC	798	GGGACUUG GCCGAAAGGCGAGUCAAGGUCU UGGGCCCA	1383
3167	CCCAGCCA G UCCUUGG	799	CCAAGGGA GCCGAAAGGCGAGUCAAGGUCU UGGCUGGG	1384
3179	CUUGGACA G CACCUUCU	800	AGAAGGUG GCCGAAAGGCGAGUCAAGGUCU UGUCCAAG	1385
3226	GGGACCUG G UGGAUGCU	801	AGCAUCCA GCCGAAAGGCGAGUCAAGGUCU CAGGUCCC	1386
3240	GCUGAGGA G UAUCUGGU	802	ACCAGAU A GCCGAAAGGCGAGUCAAGGUCU UCCUCAGC	1387
3247	AGUAUCUG G UACCCAG	803	CUGGGGUA GCCGAAAGGCGAGUCAAGGUCU CAGAUACU	1388
3255	GUACCCCA G CAGGGCUU	804	AAGCCUG GCCGAAAGGCGAGUCAAGGUCU UGGGGUAC	1389
3260	CCAGCAGG G CUUCUUCU	805	AGAAGAAG GCCGAAAGGCGAGUCAAGGUCU CUUGCUGG	1390
3287	UGCCCCGG G CGCUGGGG	806	CCCCAGCG GCCGAAAGGCGAGUCAAGGUCU CCGGGGCA	1391
3296	CGCUGGGG G CAUGGUCC	807	GGACCAUG GCCGAAAGGCGAGUCAAGGUCU CCCCAGCG	1392
3301	GGGGCAUG G UCCACCAC	808	GUGGUGGA GCCGAAAGGCGAGUCAAGGUCU CAUGCCCC	1393
3312	CACCACAG G CACCGCAG	809	CUGCGGUG GCCGAAAGGCGAGUCAAGGUCU CUGUGGUG	1394
3320	GCACCGCA G CUCAUCUA	810	UAGAUGAG GCCGAAAGGCGAGUCAAGGUCU UGCGGUGC	1395
3335	UACCAGGA G UGGCGGUG	811	CACCGCCA GCCGAAAGGCGAGUCAAGGUCU UCCUGGUA	1396
3338	CAGGAGUG G CGGUGGGG	812	CCCCACCG GCCGAAAGGCGAGUCAAGGUCU CACUCCUG	1397
3341	GAGUGGCG G UGGGGACC	813	GGUCCCA GCCGAAAGGCGAGUCAAGGUCU CGCCACUC	1398
3360	ACACUAGG G CUGGAGCC	814	GGCUCCAG GCCGAAAGGCGAGUCAAGGUCU CCUAGUGU	1399
3366	GGGUGGA G CCCUCUGA	815	UCAGAGGG GCCGAAAGGCGAGUCAAGGUCU UCCAGCCC	1400
3382	AAGAGGAG G CCCCCAGG	816	CCUGGGGG GCCGAAAGGCGAGUCAAGGUCU CUCCUCU	1401
3390	GCCCCCAG G UCUCCACU	817	AGUGGAGA GCCGAAAGGCGAGUCAAGGUCU CUGGGGGC	1402
3400	CUCCACUG G CACCCUCC	818	GGAGGGUG GCCGAAAGGCGAGUCAAGGUCU CAGUGGAG	1403
3415	CCGAAGGG G CUGGCUCC	819	GGAGCCAG GCCGAAAGGCGAGUCAAGGUCU CCCUUCGG	1404
3419	AGGGGCG G CUCCGAUG	820	CAUCGGAG GCCGAAAGGCGAGUCAAGGUCU CAGCCCCU	1405
3437	AUUUGAUG G UGACCUUG	821	CCAGGUCA GCCGAAAGGCGAGUCAAGGUCU CAUCAAU	1406
3454	GAAUGGGG G CAGCCAAG	822	CUUGGCUG GCCGAAAGGCGAGUCAAGGUCU CCCC AUUC	1407

Table 59

3457	UGGGGGCA G CCAAGGGG	823	CCCCUUGG GCCGAAAGGCGAGUCAAGGUCU	UGCCCCCA	1408
3465	GCCAAGGG G CUGCAAAG	824	CUUUGCAG GCCGAAAGGCGAGUCAAGGUCU	CCCUUGGC	1409
3473	GCUGCAAA G CCUCCCCA	825	UGGGGAGG GCCGAAAGGCGAGUCAAGGUCU	UUUGCAGC	1410
3494	UGACCCCA G CCCUCUAC	826	GUAGAGGG GCCGAAAGGCGAGUCAAGGUCU	UGGGGUCA	1411
3504	CCUCUACA G CGGUACAG	827	CUGUACCG GCCGAAAGGCGAGUCAAGGUCU	UGUAGAGG	1412
3507	CUACAGCG G UACAGUGA	828	UCACUGUA GCCGAAAGGCGAGUCAAGGUCU	CGCUGUAG	1413
3512	GCGGUACA G UGAGGACC	829	GGUCCUCA GCCGAAAGGCGAGUCAAGGUCU	UGUACCGC	1414
3526	ACCCACA G UACCCUCG	830	CAGGGGUA GCCGAAAGGCGAGUCAAGGUCU	UGUGGGGU	1415
3551	GACUGAUG G CUACGUUG	831	CAACGUAG GCCGAAAGGCGAGUCAAGGUCU	CAUCAGUC	1416
3556	AUGGCUAC G UUGCCCCC	832	GGGGGCAA GCCGAAAGGCGAGUCAAGGUCU	GUAGCCAU	1417
3575	GACCUGCA G CCCCAGC	833	GCUGGGGG GCCGAAAGGCGAGUCAAGGUCU	UGCAGGUC	1418
3582	AGCCCCCA G CCUGAAUA	834	UAUUCAGG GCCGAAAGGCGAGUCAAGGUCU	UGGGGGCU	1419
3600	GUGAACCA G CCAGAUGU	835	ACAUCUGG GCCGAAAGGCGAGUCAAGGUCU	UGGUUCAC	1420
3612	GAUGUUCG G CCCCAGCC	836	GGCUGGGG GCCGAAAGGCGAGUCAAGGUCU	CGAACAU	1421
3618	CGGCCCCA G CCCCUCU	837	GAAGGGGG GCCGAAAGGCGAGUCAAGGUCU	UGGGGGCG	1422
3638	CCGAGAGG G CCCUCUGC	838	GCAGAGGG GCCGAAAGGCGAGUCAAGGUCU	CCUCUCGG	1423
3665	ACCUCUG G UGCCACUC	839	GAGUGGCA GCCGAAAGGCGAGUCAAGGUCU	CAGCAGGU	1424
3681	CUGGAAAG G CCAAGAC	840	GUCUUGGG GCCGAAAGGCGAGUCAAGGUCU	CUUCCAG	1425
3712	AGAAUGGG G UCGUAAA	841	UUUACGA GCCGAAAGGCGAGUCAAGGUCU	CCCAUUCU	1426
3715	AUGGGGUC G UCAAAGAC	842	GUUUUGA GCCGAAAGGCGAGUCAAGGUCU	GACCCCAU	1427
3724	UCAAAGAC G UUUUUGCC	843	GGCAAAA GCCGAAAGGCGAGUCAAGGUCU	GUCUUUGA	1428
3740	CUUUGGGG G UGCCGUGG	844	CCACGGCA GCCGAAAGGCGAGUCAAGGUCU	CCCCAAAG	1429
3745	GGGUGGCC G UGAGAAC	845	GUUCUCCA GCCGAAAGGCGAGUCAAGGUCU	GGCACCCC	1430
3759	AACCCCGA G UACUUGAC	846	GUCAAGUA GCCGAAAGGCGAGUCAAGGUCU	UCGGGGUU	1431
3781	AGGGAGGA G CUGCCCCU	847	AGGGGCAG GCCGAAAGGCGAGUCAAGGUCU	UCCUCCCU	1432
3792	GCCCCUCA G CCCACCC	848	GGGUGGGG GCCGAAAGGCGAGUCAAGGUCU	UGAGGGGC	1433
3815	UGCCUUCA G CCCAGCCU	849	AGGCUGGG GCCGAAAGGCGAGUCAAGGUCU	UGAAGGCA	1434
3820	UCAGCCCA G CCUUCGAC	850	GUCGAAGG GCCGAAAGGCGAGUCAAGGUCU	UGGGCUGA	1435
3861	CCACCAGA G CGGGGGG	851	GCCCCCG GCCGAAAGGCGAGUCAAGGUCU	UCUGGUGG	1436
3868	AGCGGGGG G CUCCACCC	852	GGGUGGAG GCCGAAAGGCGAGUCAAGGUCU	CCCCCGCU	1437
3878	UCCACCCA G CACCUUCA	853	UGAAGGUG GCCGAAAGGCGAGUCAAGGUCU	UGGGUGGA	1438
3901	CACCUACG G CAGAGAAC	854	GUUCUCUG GCCGAAAGGCGAGUCAAGGUCU	CGUAGGUG	1439
3915	AACCCAGA G UACCUUGG	855	CCCAGGUA GCCGAAAGGCGAGUCAAGGUCU	UCUGGGUU	1440
3923	GUACCUUG G UCUGGACG	856	CGUCCAGA GCCGAAAGGCGAGUCAAGGUCU	CCAGGUAC	1441
3931	GUCUGGAC G UGCCAGUG	857	CACUGGCA GCCGAAAGGCGAGUCAAGGUCU	GUCCAGAC	1442
3937	ACGUGCCA G UGUAACC	858	GGUUCACA GCCGAAAGGCGAGUCAAGGUCU	UGGCACGU	1443
3951	ACCAGAAG G CCAAGUCC	859	GGACUUGG GCCGAAAGGCGAGUCAAGGUCU	CUUCUGGU	1444
3956	AAGGCCAA G UCCGCAGA	860	UCUGCGGA GCCGAAAGGCGAGUCAAGGUCU	UUGGCCUU	1445
3966	CCGCAGAA G CCCUGAUG	861	CAUCAGGG GCCGAAAGGCGAGUCAAGGUCU	UUCUGCGG	1446
3987	CUCAGGGA G CAGGAAG	862	CUUCCUG GCCGAAAGGCGAGUCAAGGUCU	UCCUGAG	1447
3996	CAGGAAG G CCUGACUU	863	AAGUCAGG GCCGAAAGGCGAGUCAAGGUCU	CUUCCUG	1448
4011	UUCUGCUG G CAUCAAGA	864	UCUUGAUG GCCGAAAGGCGAGUCAAGGUCU	CAGCAGAA	1449
4021	AUCAAGAG G UGGAGGG	865	CCCUCCA GCCGAAAGGCGAGUCAAGGUCU	CUCUUGAU	1450
4029	GUGGGAGG G CCUCCGA	866	UCGAGGG GCCGAAAGGCGAGUCAAGGUCU	CCUCCAC	1451
4100	CUGCUUGA G UUCCAGA	867	UCUGGGA GCCGAAAGGCGAGUCAAGGUCU	UCAAGCAG	1452
4111	CCCAGAUG G CUGGAAGG	868	CCUCCAG GCCGAAAGGCGAGUCAAGGUCU	CAUCUGGG	1453
4121	UGGAAGGG G UCCAGCCU	869	AGGCUUGA GCCGAAAGGCGAGUCAAGGUCU	CCCUCCA	1454

Table 59

4126	GGGGUCCA G CCUCGUUG	870	CAACGAGG GCCGAAAGGCGAGUCAAGGUCU UGGACCCC	1455
4131	CCAGCCUC G UUGGAAGA	871	UCUUCCAA GCCGAAAGGCGAGUCAAGGUCU GAGGCUGG	1456
4146	GAGGAACA G CACUGGGG	872	CCCCAGUG GCCGAAAGGCGAGUCAAGGUCU UGUUCCUC	1457
4156	ACUGGGGA G UCUUUGUG	873	CACAAAGA GCCGAAAGGCGAGUCAAGGUCU UCCCCAGU	1458
4174	AUUCUGAG G CCCUGCCC	874	GGGCAGGG GCCGAAAGGCGAGUCAAGGUCU CUCAGAAU	1459
4197	ACUCUAGG G UCCAGUGG	875	CCACUGGA GCCGAAAGGCGAGUCAAGGUCU CCUAGAGU	1460
4202	AGGGUCCA G UGGAUGCC	876	GGCAUCCA GCCGAAAGGCGAGUCAAGGUCU UGGACCCU	1461
4214	AUGCCACA G CCCAGCUU	877	AAGCUGGG GCCGAAAGGCGAGUCAAGGUCU UGUGGAU	1462
4219	ACAGCCCA G CUUGGCC	878	GGGCCAAG GCCGAAAGGCGAGUCAAGGUCU UGGGCUGU	1463
4224	CCAGCUUG G CCCUUCC	879	GGAAAGGG GCCGAAAGGCGAGUCAAGGUCU CAAGCUGG	1464
4246	GAUCCUGG G UACUGAAA	880	UUUCAGUA GCCGAAAGGCGAGUCAAGGUCU CCAGGAUC	1465
4255	UACUGAAA G CCUAGGG	881	CCCUAAGG GCCGAAAGGCGAGUCAAGGUCU UUUCAGUA	1466
4266	UUAGGGAA G CUGGCCUG	882	CAGGCCAG GCCGAAAGGCGAGUCAAGGUCU UUCCUAA	1467
4270	GGAAAGCUG G CCUGAGAG	883	CUCUCAGG GCCGAAAGGCGAGUCAAGGUCU CAGCUUCC	1468
4284	GAGGGGAA G OGGCCCUA	884	UAGGGCCG GCCGAAAGGCGAGUCAAGGUCU UUCCUC	1469
4287	GGGAAGCG G CCCUAAGG	885	CCUAGGG GCCGAAAGGCGAGUCAAGGUCU CGCUUCC	1470
4298	CUAAGGGA G UGUCUAG	886	CUUAGACA GCCGAAAGGCGAGUCAAGGUCU UCCCUUAG	1471
4314	GAACAAAA G CGACCAU	887	AUGGGUCG GCCGAAAGGCGAGUCAAGGUCU UUUUGUUC	1472
4346	GAAACCUA G UACUGCCC	888	GGGCAGUA GCCGAAAGGCGAGUCAAGGUCU UAGGUUUC	1473
4372	AAGGAACA G CAAUGGUG	889	CACCAUUG GCCGAAAGGCGAGUCAAGGUCU UGUUCCU	1474
4378	CAGCAUG G UGUCAGUA	890	UACUGACA GCCGAAAGGCGAGUCAAGGUCU CAUUGCUG	1475
4384	UGGUGUCA G UAUCCAGG	891	CCUGGAUA GCCGAAAGGCGAGUCAAGGUCU UGACACCA	1476
4392	GUAUCCAG G CUUUGUAC	892	GUACAAAG GCCGAAAGGCGAGUCAAGGUCU CUGGAUAC	1477
4404	UGUACAGA G UGCUUUC	893	GAAAAGCA GCCGAAAGGCGAGUCAAGGUCU UCUGUACA	1478
4419	UCUGUUUA G UUUUACU	894	AGUAAAAA GCCGAAAGGCGAGUCAAGGUCU UAAACAGA	1479

Input Sequence = HSERB2R. Cut Site = G/Y

Stem Length = 8 . Core Sequence = GCcgaagGCGaGuCaaGGuCu

HSERB2R (Human c-erb-B-2 mRNA; 4473 bp)

Table 60

Table 60: Substrate Specificity for Class I Ribozymes

Substrate sequence	1-9t mutation	k_{rel}
5'-GCCGU G GGUUGCAC ACCUUUCC-3'	w.t.	1.00
5'-GCCGU G GGUUGCAC ACCUUUCC-3'	A57G	2.5
5'-GCCGA G GGUUGCAC ACCUUUCC-3'	A57U	0.24
5'-GCCGC G GGUUGCAC ACCUUUCC-3'	A57G	0.66
5'-GCCGG G GGUUGCAC ACCUUUCC-3'	A57C	0.57
5'-GCCGU U GGUUGCAC ACCUUUCC-3'	w.t.	0.17
5'-GCCGU A GGUUGCAC ACCUUUCC-3'	w.t.	n.d.
5'-GCCGU C GGUUGCAC ACCUUUCC-3'	w.t.	n.d.
5'-GCCGU G GGUUGCAC ACCUUUCC-3'	C16U	0.98
5'-GCCGU G UGUUGCAC ACCUUUCC-3'	C16G	n.d.
5'-GCCGU G UGUUGCAC ACCUUUCC-3'	C16A	0.65
5'-GCCGU G AGUUGCAC ACCUUUCC-3'	C16U	0.45
5'-GCCGU G CGUUGCAC ACCUUUCC-3'	C16G	0.73
5'-GCCGU G GGUUGCAC ACCUUU-3'	w.t.	0.89
5'-GCCGU G GGUUGCAC ACCU-3'	w.t.	1.0
5'-GCCGU G GGUUGCAC AC-3'	w.t.	0.67

Table 61

Table 61: Random region alignments/mutations for Class I ribozyme

[illegible]

Table 62

Table 62: Human Her2 Class II Ribozyme and Target Sequence

RPI#	NT Pos	Substrate	Seq ID #	Ribozyme Sequence	Seq ID #
19952	433	GCUCAC G CUCACAA	7	U ₅ U ₅ G ₅ U ₅ Gag gccgaaaggGgagugagguG ₅ u gaugagc B	1480
19953	433	GCUCAC G CUCACAA	7	U ₅ U ₅ G ₅ U ₅ Gag gccgaaaggGgagugagguG ₅ u gaugagc B	1481
19950	934	CUGCCUG G CCUGCCU	22	a ₅ G ₅ G ₅ C ₅ agg gccgaaaggGgagugagguG ₅ u caggcag B	1482
19951	934	CUGCCUG G CCUGCCU	22	a ₅ G ₅ G ₅ C ₅ agg gccgaaaggGgagugagguG ₅ u caggcag B	1483
19729	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug gccgaaaggGgagugagguG ₅ u agcuca B	1484
19730	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug gccgaaaggGgagugagguG ₅ u agcuca B	1485
19731	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug gccgaaaggGgagugagguG ₅ u agcuca B	1486
20315	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ uag gccgaaaggGgagugagguG ₅ u agcucaug B	1487
20668	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ uu uaa ggc cga aag gGgagu gaG GuG ₅ uag cuc aug uuB	1488
20695	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ U ₅ U ₅ uaa agg cag aaa gG ₅ gag uga GGu G ₅ u cau guu uB	1489
20696	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ U ₅ U ₅ uaa aaggcc gaa aggGgagug aGG uG ₅ u agc uca uga uu B	1490
20719	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug gccgaaaggGgagugagguG ₅ u agcuca B	1491
20720	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug gcc P ggGgagugaGguG ₅ u agcuca B	1492
20721	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug gc P gGgagugaGguG ₅ u agcuca B	1493
20770	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ U ₅ U ₅ a ₅ ag gcc gaa agg Gga gug aGG uG ₅ u agc uca uga uu B	1494
20771	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ U ₅ U ₅ a ₅ ag gcc gaa agg Gga gug aGG uG ₅ u agc uca uga uu B	1495
20868	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug gccguuaggGgagugaGguG ₅ u agcuca B	1496
20869	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug GccgaaaggGgagugaGguG ₅ u agcuca B	1497
20870	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug GccgaaaggGgagugaGguG ₅ u agcuca B	1498
20871	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug GccgaaaggGgagugaGguG ₅ u agcuca B	1499
20872	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug gccgaaaggGgagugaGguG ₅ u agcuca B	1500
20873	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug gccgaaaggGgagugaGguG ₅ u agcuca B	1501
20874	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug gccgaaaggGgagugaGguG ₅ u agcuca B	1502
20875	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug gccgaaaggGgagugaGguG ₅ u agcuca B	1503
21448	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug g cacc GgagugaGguG ₅ u agcuca B	1504
21449	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug g uuuu GgagugaGguG ₅ u agcuca B	1505
21450	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug g uuua GgagugaGguG ₅ u agcuca B	1506
21451	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug g ucca GgagugaGguG ₅ u agcuca B	1507
21452	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug g ucuu GgagugaGguG ₅ u agcuca B	1508
21453	972	UGAGCU G CACUGC	27	G ₅ C ₅ a ₅ G ₅ ug g guua GgagugaGguG ₅ u agcuca B	1509

Table 62

21454	972	UGAGCU G CACUGC	27	9 _S C _S A _S 9 _S UG g aaU CgagugaGGuCu agcuca B	1510
21455	972	UGAGCU G CACUGC	27	9 _S C _S A _S 9 _S UG g aag CgagugaGGuCu agcuca B	1511
21456	972	UGAGCU G CACUGC	27	9 _S C _S A _S 9 _S UG g c aag g CgagugaGGuCu agcuca B	1512
21457	972	UGAGCU G CACUGC	27	9 _S C _S A _S 9 _S UG g cc aag gg CgagugaGGuCu agcuca B	1513
21458	972	UGAGCU G CACUGC	27	9 _S C _S A _S 9 _S UG g cc guua gg CgagugaGGuCu agcuca B	1514
21459	972	UGAGCU G CACUGC	27	9 _S C _S A _S 9 _S UG g cc guua gg C agugaGGuCu agcuca B	1515
19954	1292	UUUGGA G CCUGGC	34	9 _S C _S A _S 9 _S 99 gcccgaaggCgagugagguCu ucccua B	1516
20628	1292	UUUGGA G CCUGGC	34	9 _S C _S A _S 9 _S 99 GccgaaggCCGaGuGaGGuCu ucccua B	1517

lower case = 2'-O-methyl
U, C = 2'-deoxy-2'-amino U, = 2'-deoxy-2'-amino C
G,A = ribo G, A
B = inverted deoxybasic
P= polyethylene glycol 18 (PEG 18) linker

Table 63

Table 63: Human PKC α NCH Ribozyme and Substrate Sequence

Pos	Substrate	Seq ID	Ribozyme	Rz Seq ID
27	GGGGGGAC C AUGGCUGA		UCAGCCAU CUGAUGAG X CGAA IUCCCCC	
28	GGGGGACC A UGGCUGAC		GUCAGCCA CUGAUGAG X CGAA IGUCCCC	
33	ACCAUGGC U GACGUUUU		AAAACGUC CUGAUGAG X CGAA ICCAUGGU	
43	ACGUUUUC C CGGGCAAC		GUUGCCCG CUGAUGAG X CGAA IAAAACGU	
44	CGUUUUCC C GGGCAACG		CGUUGCCC CUGAUGAG X CGAA IGAAAACG	
49	UCCCGGGC A ACGACUCC		GGAGUCGU CUGAUGAG X CGAA ICCCGGGA	
55	GCAACGAC U CCACGGCG		CGCCGUGG CUGAUGAG X CGAA IUCGUUGC	
57	AACGACUC C ACGGCGUC		GACGCCGU CUGAUGAG X CGAA IAGUCGUU	
58	ACGACUCC A CGGCGUCU		AGACGCCG CUGAUGAG X CGAA IGAGUCGU	
66	ACGGCGUC U CAGGACGU		ACGUCCUG CUGAUGAG X CGAA IACGCCGU	
68	GGCGUCUC A GGACGUGG		CCACGUCC CUGAUGAG X CGAA IAGACGCC	
78	GACGUGGC C AACCGCUU		AAGCGGUU CUGAUGAG X CGAA ICCACGUC	
79	ACGUGGCC A ACGCUUC		GAAGCGGU CUGAUGAG X CGAA IGCCACGU	
82	UGGCCAAC C GCUUCGCC		GGCGAAGC CUGAUGAG X CGAA IUUGGCCA	
85	CCAACCGC U UGCCCCGC		GCGGGCGA CUGAUGAG X CGAA ICGGUUGG	
90	CGCUUCGC C CGCAAAGG		CCUUUGCG CUGAUGAG X CGAA ICGAAGCG	
91	GCUUCGCC C GCAAAGGG		CCCUUUGC CUGAUGAG X CGAA ICGAAGC	
94	UCGCCCCG A AAGGGGCG		CGCCCCUU CUGAUGAG X CGAA ICGGGCGA	
104	AGGGGCGC U GAGGCAGA		UCUGCCUC CUGAUGAG X CGAA ICGCCCCU	
110	GCUGAGGC A GAAGAACG		CGUUCUUC CUGAUGAG X CGAA ICCUCAGC	
122	GAACGUGC A CGAGGUGA		UCACCUCG CUGAUGAG X CGAA ICACGUUC	
136	UGAAGGAC C ACAAUUC		GAAUUUGU CUGAUGAG X CGAA IUCCUUCA	
137	GAAGGACC A CAAAUUCA		UGAAUUUG CUGAUGAG X CGAA IGUCCUUC	
139	AGGACCAC A AAUUCAUC		GAUGAAUU CUGAUGAG X CGAA IUUGGUCCU	
145	ACAAAUUC A UCGCGCGC		GCGCGCGA CUGAUGAG X CGAA IAAUUUGU	
154	UCGCGCGC U UCUUCAAG		CUUGAAGA CUGAUGAG X CGAA ICGCGCGA	
157	CGCGCUUC U UCAAGCAG		CUGCUUGA CUGAUGAG X CGAA IAAGCGCG	
160	GCUUCUUC A AGCAGCCC		GGGCUGCU CUGAUGAG X CGAA IAAGAAGC	
164	CUUCAAGC A GCCACCUU		AGGUGGGC CUGAUGAG X CGAA ICUUGAAG	
167	CAAGCAGC C CACCUUCU		AGAAGGUG CUGAUGAG X CGAA ICUGCUUG	
168	AAGCAGCC C ACCUUCUG		CAGAAGGU CUGAUGAG X CGAA IGCUGCUU	
169	AGCAGCCC A CCUUCUGC		GCAGAAGG CUGAUGAG X CGAA IGGCUGCU	
171	CAGCCCAC C UUCUGCAG		CUGCAGAA CUGAUGAG X CGAA IUGGGCUG	
172	AGCCCACC U UCUGCAGC		GCUGCAGA CUGAUGAG X CGAA IGUGGGCU	
175	CCACCUUC U GCAGCCAC		GUGGCUGC CUGAUGAG X CGAA IAAGGUGG	
178	CCUUCUGC A GCCACUGC		GCAGUGGC CUGAUGAG X CGAA ICAGAAGG	
181	UCUGCAGC C ACUGCACC		GGUGCAGU CUGAUGAG X CGAA ICUGCAGA	
182	CUGCAGCC A CUGCACCG		CGGUGCAG CUGAUGAG X CGAA IGCUGCAG	
184	GCAGCCAC U GCACCGAC		GUCGGUGC CUGAUGAG X CGAA IUGGCUGC	
187	GCCACUGC A CCGACUUC		GAAGUCGG CUGAUGAG X CGAA ICAGUGGC	
189	CACUGCAC C GACUUCAU		AUGAAGUC CUGAUGAG X CGAA IUGCAGUG	
193	GCACCGAC U UCAUCUGG		CCAGAUGA CUGAUGAG X CGAA IUCGGUGC	
196	CCGACUUC A UCUGGGGG		CCCCCAGA CUGAUGAG X CGAA IAAGUCGG	
199	ACUUCAUC U GGGGUVUU		AAACCCCC CUGAUGAG X CGAA IAUGAAGU	
215	UGGGAAC A AGGCUUCC		GGAAGCCU CUGAUGAG X CGAA IUUUCCTA	
220	AACAAGGC U UCCAGUGC		GCACUGGA CUGAUGAG X CGAA ICCUUGUU	
223	AAGGCUUC C AGUGCCAA		UUGGCACU CUGAUGAG X CGAA IAAGCCUU	

Table 63

224	AGGCUUCC A GUGCCAAG		CUUGGCAC CUGAUGAG X CGAA IGAAGCCU	
229	UCCAGUGC C AAGUUUGC		GCAAACU CUGAUGAG X CGAA ICACUGGA	
230	CCAGUGCC A AGUUUGCU		AGCAAACU CUGAUGAG X CGAA IGCACUGG	
238	AAGUUUGC U GUUUUGUG		CACAAAAC CUGAUGAG X CGAA ICAAACUU	
250	UUGUGGUC C ACAAGAGG		CCUCUUGU CUGAUGAG X CGAA IACCACAA	
251	UGUGGUCC A CAAGAGGU		ACCUCUUG CUGAUGAG X CGAA IGACCACA	
253	UGGUCCAC A AGAGGUGC		GCACCUCU CUGAUGAG X CGAA IUGGACCA	
262	AGAGGUGC C AUGAAUUU		AAAUUCAU CUGAUGAG X CGAA ICACCUCU	
263	GAGGUGCC A UGAAUUUG		CAAAUUCA CUGAUGAG X CGAA IGCACCUC	
276	UUUGUUAC U UUUUCUUG		CAAGAAAA CUGAUGAG X CGAA IUAACAAA	
282	ACUUUUUC U UGUCCGGG		CCCGGACA CUGAUGAG X CGAA IAAAAAGU	
287	UUCUUGUC C GGGUGCGG		CCGCACCC CUGAUGAG X CGAA IACAAGAA	
305	UAAGGGAC C CGACACUG		CAGUGUCG CUGAUGAG X CGAA IUCCCUUA	
306	AAGGGACC C GACACUGA		UCAGUGUC CUGAUGAG X CGAA IGUCCCUU	
310	GACCCGAC A CUGAUGAC		GUCAUCAG CUGAUGAG X CGAA IUCCGGUC	
312	CCCGACAC U GAUGACCC		GGGUCAUC CUGAUGAG X CGAA IUGUCGGG	
319	CUGAUGAC C CCAGGAGC		GCUCCUGG CUGAUGAG X CGAA IUCAUCAG	
320	UGAUGACC C CAGGAGCA		UGCUCUG CUGAUGAG X CGAA IGUCAUCA	
321	GAUGACCC C AGGAGCAA		UUGCUCU CUGAUGAG X CGAA IGGUCAUC	
322	AUGACCCC A GGAGCAAG		CUUGCUC CUGAUGAG X CGAA IGGUCAU	
328	CCAGGAGC A AGCACAAG		CUUGUGCU CUGAUGAG X CGAA ICUCCUGG	
332	GAGCAAGC A CAAGUUCA		UGAACTUG CUGAUGAG X CGAA ICUUGCUC	
334	GCAAGCAC A AGUUCAAA		UUUGAACU CUGAUGAG X CGAA IUGCUUGC	
340	ACAAGUUC A AAUCCAC		GUGGAUUU CUGAUGAG X CGAA IAACUUGU	
346	UCAAAUUC C ACACUAC		GUAAGUGU CUGAUGAG X CGAA IAUUUUGA	
347	CAAAAUCC A CACUACG		CGUAAGUG CUGAUGAG X CGAA IGAUUUUG	
349	AAAUCCAC A CUACGGA		UCCGUUAG CUGAUGAG X CGAA IUGGAUUU	
351	AUCCACAC U UACGGAAG		CUUCCGUA CUGAUGAG X CGAA IUGUGGAU	
361	ACGGAAGC C CCACCUUC		GAAGGUGG CUGAUGAG X CGAA ICUUCCGU	
362	CGGAAGCC C CACCUUCU		AGAAGGUG CUGAUGAG X CGAA IGCUCCG	
363	GGAAGCCC C ACCUUCUG		CAGAAGGU CUGAUGAG X CGAA IGGCUUCC	
364	GAAGCCCC A CCUUCUGC		GCAGAAGG CUGAUGAG X CGAA IGGCUUC	
366	AGCCCCAC C UUCUGCGA		UCGCAGAA CUGAUGAG X CGAA IUGGGGCU	
367	GCCCCACC U UCUGCGAU		AUCGCAGA CUGAUGAG X CGAA IGUGGGGC	
370	CCACCUUC U GCGAUCAC		GUGAUCGC CUGAUGAG X CGAA IAAGGUGG	
377	CUGCGAUC A CUGUGGGU		ACCCACAG CUGAUGAG X CGAA IAUCGCAG	
379	GCGAUCAC U GUGGGUCA		UGACCCAC CUGAUGAG X CGAA IUGAUCGC	
387	UGUGGGUC A CUGCUCUA		UAGAGCAG CUGAUGAG X CGAA IACCCACA	
389	UGGUCAC U GCUCUAUG		CAUAGAGC CUGAUGAG X CGAA IUGACCCA	
392	GUCACUGC U CUAUGGAC		GUCCAUG CUGAUGAG X CGAA ICAGUGAC	
394	CACUGCUC U AUGGACUU		AAGUCCA CUGAUGAG X CGAA IAGCAGUG	
401	CUAUGGAC U UAUCCAUC		GAUGGAU CUGAUGAG X CGAA IUCCAUG	
406	GACUUAUC C AUCAAGGG		CCCUUGAU CUGAUGAG X CGAA IAUAGUC	
407	ACUUAUCC A UCAAGGGA		UCCCUUGA CUGAUGAG X CGAA IGAUAAGU	
410	UAUCCAUC A AGGGAUGA		UCAUCCU CUGAUGAG X CGAA IAUGGAUA	
427	AAUGUGAC A CCUGCGAU		AUCGCAGG CUGAUGAG X CGAA IUCACAUU	
429	UGUGACAC C UGCGAUAU		AUAUCGCA CUGAUGAG X CGAA IUGUCACA	
430	GUGACACC U GCGAU AUG		CAUAUCGC CUGAUGAG X CGAA IGUGUCAC	
446	GAACGUUC A CAAGCAAU		AUUGCUUG CUGAUGAG X CGAA IAACGUUC	
448	ACGUUCAC A AGCAAUGC		GCAUUGCU CUGAUGAG X CGAA IUGAACGU	
452	UCACAAGC A AUGCGUCA		UGACGCAU CUGAUGAG X CGAA ICUUGUGA	

Table 63

460	AAUGCGUC A UCAAUGUC		GACAUUGA CUGAUGAG X CGAA IACGCAUJ	
463	GCGUCAUC A AUGUCCCC		GGGGACAU CUGAUGAG X CGAA IAUGACGC	
469	UCAAUGUC C CCAGCCUC		GAGGCUGG CUGAUGAG X CGAA IACAUUGA	
470	CAAUGUCC C CAGCCUCU		AGAGGCUG CUGAUGAG X CGAA IGACAUUG	
471	AAUGUCCC C AGCCUCUG		CAGAGGCU CUGAUGAG X CGAA IGGACAUU	
472	AUGUCCCC A GCCUCUGC		GCAGAGGC CUGAUGAG X CGAA IGGGACAU	
475	UCCCCAGC C UCUGCGGA		UCCGCAGA CUGAUGAG X CGAA ICUGGGGA	
476	CCCCAGCC U CUGCGGAA		UCCGCAG CUGAUGAG X CGAA IGCUGGGG	
478	CCAGCCUC U GCGGAAUG		CAUCCGC CUGAUGAG X CGAA IAGGCUGG	
491	AAUGGAUC A CACUGAGA		UCUCAGUG CUGAUGAG X CGAA IAUCCAUI	
493	UGGAUCAC A CUGAGAAG		CUUCUCAG CUGAUGAG X CGAA IUGAUCCA	
495	GAUCACAC U GAGAAGAG		CUCUUCUC CUGAUGAG X CGAA IUGUGAUC	
517	GGAUUUAC C UAAAGGCU		AGCCUUUA CUGAUGAG X CGAA IUAAAUCC	
518	GAUUUACC U AAAGGCUG		CAGCCUUU CUGAUGAG X CGAA IGUAAAUC	
525	CUAAAGGC U GAGGUUGC		GCAACCUC CUGAUGAG X CGAA ICCUUUAG	
534	GAGGUUGC U GAUGAAAA		UUUUCAUC CUGAUGAG X CGAA ICAACCUC	
545	UGAAAAGC U CCAUGUCA		UGACAUGG CUGAUGAG X CGAA ICUUUUCA	
547	AAAAGCUC C AUGUCACA		UGUGACAU CUGAUGAG X CGAA IAGCUUUU	
548	AAAGCUCC A UGUCACAG		CUGUGACA CUGAUGAG X CGAA IGAGCUUU	
553	UCCAUGUC A CAGUACGA		UCGUACUG CUGAUGAG X CGAA IACAUGGA	
555	CAUGUCAC A GUACGAGA		UCUCGUAC CUGAUGAG X CGAA IUGACAUG	
567	CGAGAUGC A AAAAAUCU		AGAUUUUU CUGAUGAG X CGAA ICAUCUCG	
575	AAAAAUC U AAUCCCUA		UAGGGAUU CUGAUGAG X CGAA IAUUUUUU	
580	AUCUAAUC C CUAUGGAU		AUCCAUA CUGAUGAG X CGAA IAUUAGAU	
581	UCUAAUCC C UAUGGAUC		GAUCCAUA CUGAUGAG X CGAA IGAUUAGA	
582	CUAAUCCC U AUGGAUCC		GGAUCCA CUGAUGAG X CGAA IGGAUUAG	
590	UAUGGAUC C AAACGGGC		GCCCGUUU CUGAUGAG X CGAA IAUCCAUA	
591	AUGGAUCC A AACGGGCU		AGCCGCUU CUGAUGAG X CGAA IGAUCCA	
599	AAACGGGC U UUCAGAUC		GAUCUGAA CUGAUGAG X CGAA ICCCGUUU	
603	GGGCUUUC A GAUCCUUA		UAAGGAUC CUGAUGAG X CGAA IAAAGCCC	
608	UUCAGAUC C UUAUGUGA		UCACAUAA CUGAUGAG X CGAA IAUUGGAA	
609	UCAGAUC U UAUGUGAA		UUCACAU CUGAUGAG X CGAA IGAUCUGA	
620	UGUGAAGC U GAAACUUA		UAAGUUUC CUGAUGAG X CGAA ICUUCACA	
626	GCUGAAAC U UAUUCCUG		CAGGAAUA CUGAUGAG X CGAA IUUUCAGC	
632	ACUUUUC C UGAUCCCA		UGGGAUCA CUGAUGAG X CGAA IAAUAAGU	
633	CUUUAUCC U GAUCCCAA		UUGGGAUC CUGAUGAG X CGAA IGAAUAAG	
638	UCCUGAUC C CAAGAAUG		CAUUCUUG CUGAUGAG X CGAA IAUCAAGG	
639	CCUGAUCC C AAGAAUGA		UCAUUCU CUGAUGAG X CGAA IGAUCAGG	
640	CUGAUCCC A AGAAUGAA		UUCAUUCU CUGAUGAG X CGAA IGGAUCA	
652	AUGAAAGC A AGCAAAAA		UUUUUGCU CUGAUGAG X CGAA ICUUUCAU	
656	AAGCAAGC A AAAAACCA		UGGUUUUU CUGAUGAG X CGAA ICUUGCUU	
663	CAAAAAAC C AAAACCAU		AUGGUUUU CUGAUGAG X CGAA IUUUUUUG	
664	AAAAAAC C A AAACCAUC		GAUGUUUU CUGAUGAG X CGAA IGUUUUUU	
669	ACCAAAAC C AUCCGCUC		GAGCGGAU CUGAUGAG X CGAA IUUUUGGU	
670	CCAAAC C AUCCGCUC		GGAGCGGA CUGAUGAG X CGAA IGUUUUGG	
673	AAACCAUC C GCUCCACA		UGUGGAGC CUGAUGAG X CGAA IAUGGUUU	
676	CCAUCGC C CCACACUA		UAGUGUGG CUGAUGAG X CGAA ICGAUGG	
678	AUCCGCUC C ACACUAAA		UUUAGUGU CUGAUGAG X CGAA IAGCGGAU	
679	UCCGCUC C ACACUAAA		AUUUAGUG CUGAUGAG X CGAA IGAGCGGA	
681	CGCUCCAC A CUAAAUCC		GGAUUUAG CUGAUGAG X CGAA IUUGAGCG	
683	CUCCACAC U AAUCCGC		GCGGAUUU CUGAUGAG X CGAA IUUGGAG	

Table 63

689	ACUAAAUC C GCAGUGGA		UCCACUGC CUGAUGAG X CGAA IAUUUAGU	
692	AAAUCCGC A GUGGAAUG		CAUUCAC CUGAUGAG X CGAA ICGAAUUU	
705	AAUGAGUC C UUUACAUU		AAUGUAAA CUGAUGAG X CGAA IACUCAU	
706	AUGAGUCC U UUACAUUC		GAAUGUAA CUGAUGAG X CGAA IGACUCAU	
711	UCCUUUAC A UUCAAAU		AAUUUGAA CUGAUGAG X CGAA IUAAAGGA	
715	UUACAUUC A AAUUGAAA		UUUCAAUU CUGAUGAG X CGAA IAAUGUAA	
725	AUUGAAAC C UUCAGACA		UGUCUGAA CUGAUGAG X CGAA IUUUCAAU	
726	UUGAAACC U UCAGACAA		UUGUCUGA CUGAUGAG X CGAA IGUUUCAA	
729	AAACCUUC A GACAAAGA		UCUUUGUC CUGAUGAG X CGAA IAAGGUUU	
733	CUUCAGAC A AAGACCGA		UCGGUCUU CUGAUGAG X CGAA IUCUGAAG	
739	ACAAAGAC C GACGACUG		CAGUCGUC CUGAUGAG X CGAA IUCUUUGU	
746	CCGACGAC U GUCUGUAG		CUACAGAC CUGAUGAG X CGAA IUCGUCGG	
750	CGACUGUC U GUAGAAAU		AUUUCUAC CUGAUGAG X CGAA IACAGUCG	
760	UAGAAAUU U GGGACUGG		CCAGUCCC CUGAUGAG X CGAA IAUUUUA	
766	UCUGGGAC U GGGAUCCA		UCGAUCCC CUGAUGAG X CGAA IUCCGAGA	
777	GAUCGAAC A ACAAGGAA		UUCCUUU CUGAUGAG X CGAA IUUCGAUC	
780	CGAACAA C A AGGAUGA		UCAUUCU CUGAUGAG X CGAA IUUGUUCG	
790	GGAUGAC U UCAUGGGA		UCCCAUGA CUGAUGAG X CGAA IUCAUUC	
793	AUGACUUC A UGGGAUCC		GGAUCCCA CUGAUGAG X CGAA IAAGUCAU	
801	AUGGGAUC C CUUUCUU		AAGGAAAG CUGAUGAG X CGAA IAUCCAU	
802	UGGGAUCC C UUUCUUU		AAAGGAAA CUGAUGAG X CGAA IGAUCCCA	
803	GGGAUCCC U UUUCUUU		CAAAGGAA CUGAUGAG X CGAA IGGAUCCC	
807	UCCCUUUC C UUUGGAGU		ACUCCAAA CUGAUGAG X CGAA IAAAGGGA	
808	CCCUUUCU U UUGGAGUU		AACUCCAA CUGAUGAG X CGAA IGAAAGGG	
824	UUCGGAGC U GAUGAAGA		UCUUCUAC CUGAUGAG X CGAA ICUCGAA	
836	GAAGAGUC C GGCCAGUG		CACUGGCC CUGAUGAG X CGAA ICAUCUUC	
840	AUGCCGGC C AGUGGAUG		CAUCCACU CUGAUGAG X CGAA ICCGGCAU	
841	UGCCGGCC A GUGGAUGG		CCAUCCAC CUGAUGAG X CGAA IGCCGGCA	
853	GAUGGUAC A AGUUGCUU		AAGCAACU CUGAUGAG X CGAA IUACCAUC	
860	CAAGUUC U UAACCAAG		CUUGGUUA CUGAUGAG X CGAA ICAACUUG	
865	UGCUAAC C AAGAAGAA		UUUUUUU CUGAUGAG X CGAA IUUAAGCA	
866	GCUUAACC A AGAAGAA		CUUCUUU CUGAUGAG X CGAA IGUAAGC	
883	GUGAGUAC U ACAACGUA		UACGUUGU CUGAUGAG X CGAA IUACUCAC	
886	AGUACUAC A ACGUACCC		GGGUACGU CUGAUGAG X CGAA IUAGUACU	
893	CAACGUAC C CAUUCGG		CCGGAAUG CUGAUGAG X CGAA IUACGUUG	
894	AACGUACC C AUUCGGAA		UCCGGAAU CUGAUGAG X CGAA IGUAAGUU	
895	ACGUACCC A UUCCGGAA		UUCCGGAA CUGAUGAG X CGAA IGGUACGU	
899	ACCAUUC C GGAAGGG		CCCUUUC CUGAUGAG X CGAA IAAUGGGU	
922	AAGGAAAC A UGGAACUC		GAGUUCCA CUGAUGAG X CGAA IUUUCCUU	
929	CAUGGAAC U CAGGCAGA		UCUGCCUG CUGAUGAG X CGAA IUUCCAUG	
931	UGGAACUC A GGCAGAAA		UUUCUGCC CUGAUGAG X CGAA IAGUCCA	
935	ACUCAGGC A GAAAUUCG		CGAAUUUC CUGAUGAG X CGAA ICCUGAGU	
951	GAGAAAGC C AAACUUGG		CCAAGUUU CUGAUGAG X CGAA ICUUUCUC	
952	AGAAAGCC A AACUUGG		GCCAAGUU CUGAUGAG X CGAA IGCUUUCU	
956	AGCCAAAC U UGGCCUG		CAGGGCCA CUGAUGAG X CGAA IUUUGGCU	
961	AACUUGGC C CUGCUGGC		GCCAGCAG CUGAUGAG X CGAA ICCAAGUU	
962	ACUUGGCC C UGCUGGCA		UGCCAGCA CUGAUGAG X CGAA IGCCAAGU	
963	CUUGGCC U GCUGGCAA		UUGCCAGC CUGAUGAG X CGAA IGGCCAAG	
966	GGCCUGC U GGCAACAA		UUGUUGCC CUGAUGAG X CGAA ICAGGGCC	
970	CUGCUGGC A ACAAAGUC		GACUUUGU CUGAUGAG X CGAA ICCAGCAG	
973	CUGGCAAC A AAGUCAUC		GAUGACUU CUGAUGAG X CGAA IUUGCCAG	

Table 63

979	ACAAAGUC A UCAGUCCC		GGGACUGA CUGAUGAG X CGAA IACUUUGU	
982	AAGUCAUC A GUCCCUCU		AGAGGGAC CUGAUGAG X CGAA IAUGACUU	
986	CAUCAGUC C CUCUGAAG		CUUCAGAG CUGAUGAG X CGAA IACUGAUG	
987	AUCAGUCC C UCUGAAGA		UCUUCAGA CUGAUGAG X CGAA IGACUGAU	
988	UCAGUCCC U CUGAAGAC		GUCUUCAG CUGAUGAG X CGAA IGGACUGA	
990	AGUCCCUC U GAAGACAG		CUGUCUUC CUGAUGAG X CGAA IAGGGACU	
997	CUGAAGAC A GGAAACAA		UUGUUUCC CUGAUGAG X CGAA IUCUUCAG	
1004	CAGGAAAC A ACCUCCA		UGGAAGGU CUGAUGAG X CGAA IUUCCUG	
1007	GAAACAAC C UCCAACA		UGUUGGAA CUGAUGAG X CGAA IUUGUUUC	
1008	AAACAACC U UCCAACA		UUGUUGGA CUGAUGAG X CGAA IGUUGUUU	
1011	CAACCUUC C AACAACCU		AGGUUGUU CUGAUGAG X CGAA IAAGGUUG	
1012	AACCUUCC A ACAACCU		AAGGUUGU CUGAUGAG X CGAA IGAAGGUU	
1015	CUUCCAAC A ACCUUGAC		GUCAAGGU CUGAUGAG X CGAA IUUGGAAG	
1018	CCAACAAC C UUGACCGA		UCGGUCAA CUGAUGAG X CGAA IUUGUUGG	
1019	CAACAACC U UGACCGAG		CUCGGUCA CUGAUGAG X CGAA IGUUGUUG	
1024	ACCUUGAC C GAGUGAAA		UUUCACUC CUGAUGAG X CGAA IUCAAGGU	
1034	AGUGAAAC U CACGGACU		AGUCCGUG CUGAUGAG X CGAA IUUUCACU	
1036	UGAAACUC A CGGACUUC		GAAGUCCG CUGAUGAG X CGAA IAGUUUCA	
1042	UCACGGAC U UCAAUUUC		GAAAUUGA CUGAUGAG X CGAA IUCCGUGA	
1045	CGGACUUC A AUUCCUC		GAGGAAAU CUGAUGAG X CGAA IAAGUCCG	
1051	UCAAUUUC C UCAUGGUG		CACCAUGA CUGAUGAG X CGAA IAAAUUGA	
1052	CAAUUUCC U CAUGGUGU		ACACCAUG CUGAUGAG X CGAA IGAAAUUG	
1054	AUUUCCUC A UGGUGUUG		CAACACCA CUGAUGAG X CGAA IAGGAAAU	
1091	GGUGAUGC U UGCCGACA		UGUCGGCA CUGAUGAG X CGAA ICAUCACC	
1095	AUGCUGUC C GACAGGAA		UUCUGUC CUGAUGAG X CGAA ICAAGCAU	
1099	UUGCCGAC A GGAAGGGC		GCCUUC CUGAUGAG X CGAA IUCGGCAA	
1108	GGAAGGGC A CAGAAGAA		UUCUUCUG CUGAUGAG X CGAA ICCCUUCC	
1110	AAGGGCAC A GAAGAAU		AGUUCUUC CUGAUGAG X CGAA IUGCCCUU	
1118	AGAAGAAC U GUAUGCAA		UUGCAUAC CUGAUGAG X CGAA IUUCUUCU	
1125	CUGUAUGC A AUCAAAAU		AUUUUGAU CUGAUGAG X CGAA ICAUACAG	
1129	AUGCAAUC A AAAUCCUG		CAGGAUUU CUGAUGAG X CGAA IAUUGCAU	
1135	UCAAAAU C UGAAGAAG		CUUCUUCA CUGAUGAG X CGAA IAUUUUGA	
1136	CAAAAUCC U GAAGAAGG		CCUUCUUC CUGAUGAG X CGAA IGAUUUUG	
1157	GGUGAUUC A GGAUGAUG		CAUCAUCC CUGAUGAG X CGAA IAAUACCC	
1177	UGGAGUGC A CCAUGGUA		UACCAUGG CUGAUGAG X CGAA ICACUCCA	
1179	GAGUGCAC C AUGGUAGA		UCUACCAU CUGAUGAG X CGAA IUGCACUC	
1180	AGUGCACC A UGGUAGAA		UUCUACCA CUGAUGAG X CGAA IGUGCACU	
1198	AGCGAGUC U UGGCCUG		CAGGGCCA CUGAUGAG X CGAA IACUCGCU	
1203	GUCUUGGC C CUGCUUGA		UCAAGCAG CUGAUGAG X CGAA ICCAAGAC	
1204	UCUUGGCC C UGCUUGAC		GUCAAGCA CUGAUGAG X CGAA IGCCAAGA	
1205	CUUGGCCC U GCUUGACA		UGUCAAGC CUGAUGAG X CGAA IGGCCAAG	
1208	GGCCUGC U UGACAAAC		GUUUGUCA CUGAUGAG X CGAA ICAGGGCC	
1213	UGCUGAC A AACCCTCG		CGGGGUU CUGAUGAG X CGAA IUCAAGCA	
1217	UGACAAAC C CCCGUUCU		AGAACGGG CUGAUGAG X CGAA IUUUGUCA	
1218	GACAAACC C CCGUUCU		AAGAACGG CUGAUGAG X CGAA IGUUGUC	
1219	ACAAACCC C CGUUCUUG		CAAGAACG CUGAUGAG X CGAA IGGUUGU	
1220	CAAACCCC C GUUCUUGA		UCAAGAAC CUGAUGAG X CGAA IGGGUUG	
1225	CCCCGUUC U UGACGCAG		CUGCGUCA CUGAUGAG X CGAA IAACGGGG	
1232	CUUGACGC A GCUGCAU		AGUGCAGC CUGAUGAG X CGAA ICGUCAAG	
1235	GACGCAGC U GCACUCCU		AGGAGUGC CUGAUGAG X CGAA ICUGCGUC	
1238	GCAGCUGC A CUCCUGCU		AGCAGGAG CUGAUGAG X CGAA ICAGCUGC	

Table 63

1240	AGCUGCAC U CCUGCUUC		GAAGCAGG CUGAUGAG X CGAA IUGCAGCU	
1242	CUGCACUC C UGCUUCCA		UGGAAGCA CUGAUGAG X CGAA IAGUGCAG	
1243	UGCACUCC U GCUUCCAG		CUGGAAGC CUGAUGAG X CGAA IGAGUGCA	
1246	ACUCCUGC U UCCAGACA		UGUCUGGA CUGAUGAG X CGAA ICAGGAGU	
1249	CCUGCUUC C AGACAGUG		CACUGUCU CUGAUGAG X CGAA IAAGCAGG	
1250	CUGCUUCC A GACAGUGG		CCACUGUC CUGAUGAG X CGAA IGAAGCAG	
1254	UUCACAGC A GUGGAUCG		CGAUCCAC CUGAUGAG X CGAA IUCUGGAA	
1265	GGAUCGGC U GUACUUCG		CGAAGUAC CUGAUGAG X CGAA ICCGAUCC	
1270	GGCUGUAC U UCGUCAUG		CAUGACGA CUGAUGAG X CGAA IUACAGCC	
1276	ACUUCGUC A UGGAUAU		AUAUCCA CUGAUGAG X CGAA IACGAAGU	
1288	AAUAUGUC A ACGGUGGG		CCCACCGU CUGAUGAG X CGAA IACAUUU	
1300	GUGGGGAC C UCAUGUAC		GUACAUGA CUGAUGAG X CGAA IUCCCCAC	
1301	UGGGGACC U CAUGUACC		GGUACAUG CUGAUGAG X CGAA IGUCCCCA	
1303	GGGACCUC A UGUACCAC		GUGGUACA CUGAUGAG X CGAA IAGGUCCC	
1309	UCAUGUAC C ACAUUCAG		CUGAAUGU CUGAUGAG X CGAA IUACAUGA	
1310	CAUGUACC A CAUUCAGC		GCUGAAUG CUGAUGAG X CGAA IGUACAUG	
1312	UGUACCAC A UUCAGCAA		UUGCUGAA CUGAUGAG X CGAA IUGGUACA	
1316	CCACAUUC A GCAAGUAG		CUACUUGC CUGAUGAG X CGAA IAAUGUGG	
1319	CAUUCAGC A AGUAGGAA		UUCCUACU CUGAUGAG X CGAA ICUGAAUG	
1340	UAAGGAAC C ACAAGCAG		CUGCUUGU CUGAUGAG X CGAA IUUCCUUA	
1341	AAGGAACC A CAAGCAGU		ACUGCUUG CUGAUGAG X CGAA IGUCCUUA	
1343	GGAACCAC A AGCAGUAU		AUACUGCU CUGAUGAG X CGAA IUGGUUCC	
1347	CCACAAGC A GUAUUCUA		UAGAAUAC CUGAUGAG X CGAA ICUUGUGG	
1354	CAGUAUUC U AUGCGGCA		UGCCGCAU CUGAUGAG X CGAA IAAUACUG	
1362	UAUGCGGC A GAGAUUUC		GAAAUUC CUGAUGAG X CGAA ICCGCAUA	
1371	GAGAUUUC C AUCGGAUU		AAUCCGAU CUGAUGAG X CGAA IAAAUCUC	
1372	AGAUUUCC A UCGGAUUG		CAAUCCGA CUGAUGAG X CGAA IGAAAUUC	
1384	GAUUGUUC U UUCUUAU		AUGAAGAA CUGAUGAG X CGAA IAACAAUC	
1388	GUUCUUUC U UCAUAAAA		UUUUAUGA CUGAUGAG X CGAA IAAAGAAC	
1391	CUUUCUUC A UAAAAGAG		CUCUUUUA CUGAUGAG X CGAA IAAGAAAG	
1405	GAGGAUUC A UUAUAGG		CCUAUAAA CUGAUGAG X CGAA IAUUCCUC	
1418	UAGGGAUC U GAAGUUAG		CUAACUUC CUGAUGAG X CGAA IAUCCCUA	
1435	AUAACGUC A UGUUGGAU		AUCCAACA CUGAUGAG X CGAA IACGUUAU	
1446	UUGGAUUC A GAAGGACA		UGUCCUUC CUGAUGAG X CGAA IAAUCCAA	
1454	AGAAGGAC A UAUCAAAA		UUUUGAUA CUGAUGAG X CGAA IUCCUUCU	
1459	GACAUUUC A AAAUUGCU		AGCAAUUU CUGAUGAG X CGAA IAUUUGUC	
1467	AAAAUUGC U GACUUUGG		CCAAAGUC CUGAUGAG X CGAA ICAUUUUU	
1471	UUGCUGAC U UUGGGAUG		CAUCCCAA CUGAUGAG X CGAA IUCAGCAA	
1483	GGAUGUGC A AGGAACAC		GUGUUCU CUGAUGAG X CGAA ICACAUCC	
1490	CAAGGAAC A CAUGAUGG		CCAUCUUG CUGAUGAG X CGAA IUUCCUUG	
1492	AGGAACAC A UGAUGGAU		AUCCAACA CUGAUGAG X CGAA IUGUCCU	
1507	AUGGAGUC A CGACCAGG		CCUGGUCG CUGAUGAG X CGAA IACUCCAU	
1512	GUCACGAC C AGGACCUU		AAGGUCCU CUGAUGAG X CGAA IUCGUGAC	
1513	UCACGACC A GGACCUUC		GAAGGUCC CUGAUGAG X CGAA IGUCGUGA	
1518	ACCAGGAC C UUCUGUGG		CCACAGAA CUGAUGAG X CGAA IUCCUGGU	
1519	CCAGGACC U UCUGUGGG		CCACAGAA CUGAUGAG X CGAA IGUCUGG	
1522	GGACCUUC U GUGGGACU		AGUCCAC CUGAUGAG X CGAA IAAGGUCC	
1530	UGUGGGAC U CCAGAUUA		UAAUCUGG CUGAUGAG X CGAA IUCCACA	
1532	UGGGACUC C AGAUUAUA		UAUAAUCU CUGAUGAG X CGAA IAGUCCCA	
1533	GGGACUCC A GAUUAUAU		AUAUAAUC CUGAUGAG X CGAA IGAGUCCC	
1545	UAUAUCGC C CCAGAGAU		AUCUCUGG CUGAUGAG X CGAA ICGAUUA	

Table 63

1546	AUAUCGCC C CAGAGUA		UAUCUCUG CUGAUGAG X CGAA IGCGAUU	
1547	UAUCGCCC C AGAGUAA		UUAUCUCU CUGAUGAG X CGAA IGGCGAU	
1548	AUCGCCCC A GAGUAAU		AUUAUCUC CUGAUGAG X CGAA IGGGCGAU	
1560	AUAAUCGC U UAUCAGCC		GGCUGAUA CUGAUGAG X CGAA ICGAUU	
1565	CGCUUAUC A GCCGUAUG		CAUACGGC CUGAUGAG X CGAA IAUAGCG	
1568	UUAUCAGC C GUAUGGAA		UUCCAUAC CUGAUGAG X CGAA ICUGAUAA	
1581	GGAAAAUC U GUGGACUG		CAGUCCAC CUGAUGAG X CGAA IAUUUCC	
1588	CUGUGGAC U GGUGGGCC		GGCCACC CUGAUGAG X CGAA IUCCACAG	
1596	UGGUGGC C UAUGGCGU		ACGCCAU CUGAUGAG X CGAA ICCACCA	
1597	GGUGGGCC U AUGGCGUC		GACGCCAU CUGAUGAG X CGAA IGGCCACC	
1606	AUGGCGUC C UGUUGUAU		AUACAACA CUGAUGAG X CGAA IACGCCAU	
1607	UGGCGUCC U GUUGUAUG		CAUACAAC CUGAUGAG X CGAA IGACGCCA	
1622	UGAAUUGC U UGCCGGGC		GCCCGGCA CUGAUGAG X CGAA ICAUUUCA	
1626	AUGCUGC C GGCAGCC		GGCUGCCC CUGAUGAG X CGAA ICAAGCAU	
1631	UGCCGGGC A GCCUCCAU		AUGGAGGC CUGAUGAG X CGAA ICCCGGCA	
1634	CGGGCAGC C UCCAUUUG		CAAAUGGA CUGAUGAG X CGAA ICUGCCC	
1635	GGGCAGCC U CCAUUUGA		UCAAUGG CUGAUGAG X CGAA IAGCUGCC	
1637	GCAGCCUC C AUUGAUG		CAUCAAAU CUGAUGAG X CGAA IAGGCUGC	
1638	CAGCCUCC A UUUGAUGG		CCAUCAAA CUGAUGAG X CGAA IAGGCUG	
1664	AGACGAGC U AUUUCAGU		ACUGAAAU CUGAUGAG X CGAA ICUCGUCU	
1670	GCUAUUC A GUCUAUCA		UGAUAGAC CUGAUGAG X CGAA IAAAUAGC	
1674	UUUCAGUC U AUCAUGGA		UCCAUGAU CUGAUGAG X CGAA IACUGAAA	
1678	AGUCUAUC A UGGAGCAC		GUGCUCCA CUGAUGAG X CGAA IAUAGACU	
1685	CAUGGAGC A CAACGUUU		AAACGUUG CUGAUGAG X CGAA ICUCCAUG	
1687	UGGAGCAC A ACGUUUCC		GGAAACGU CUGAUGAG X CGAA IUGCUCCA	
1695	AACGUUUC C UAUCCAAA		UUUGGAUA CUGAUGAG X CGAA IAAACGUU	
1696	ACGUUUC C AUCCAAA		UUUUGGAU CUGAUGAG X CGAA IGAAACGU	
1700	UUCUAUC C AAAUCCU		AGGAUUUU CUGAUGAG X CGAA IAUAGGAA	
1701	UCCUAUC C AAAUCCU		AAGGAUUU CUGAUGAG X CGAA IGAUAGGA	
1707	CCAAAUC C UUGUCCAA		UUGGACAA CUGAUGAG X CGAA IAUUUUGG	
1708	CAAAUCC U UGUCCAAG		CUUGGACA CUGAUGAG X CGAA IGAUUUG	
1713	UCCUUGUC C AAGGAGGC		GCCUCCU CUGAUGAG X CGAA IACAAGGA	
1714	CCUUGUCC A AGGAGGCU		AGCCUCCU CUGAUGAG X CGAA IGACAAGG	
1722	AAGGAGGC U GUUUCUAU		AUAGAAAC CUGAUGAG X CGAA ICCUCCU	
1728	GCUGUUC U AUCUGCAA		UUGCAGAU CUGAUGAG X CGAA IAAACAGC	
1732	UUUCUAUC U GCAAAGGA		UCCUUGC CUGAUGAG X CGAA IAUAGAAA	
1735	CUAUCUGC A AAGGACUG		CAGUCCU CUGAUGAG X CGAA ICAGAUAG	
1742	CAAAGGAC U GAUGACCA		UGGUCAUC CUGAUGAG X CGAA IUCCUUG	
1749	CUGAUGAC C AAACACCC		GGUGUUU CUGAUGAG X CGAA IUCAUCAG	
1750	UGAUGACC A AACACCCA		UGGGUGU CUGAUGAG X CGAA IGUCAUCA	
1754	GACCAAAC A CCCAGCCA		UGGCGGG CUGAUGAG X CGAA IUUUGGUC	
1756	CCAAACAC C CAGCCAAG		CUUGGCUG CUGAUGAG X CGAA IUGUUUGG	
1757	CAAACACC C AGCCAAGC		GCUUGGCU CUGAUGAG X CGAA IGUGUUUG	
1758	AAACACCC A GCCAAGCG		CGCUUGGC CUGAUGAG X CGAA IGGUGUUU	
1761	CACCCAGC C AAGCGGCU		AGCCGCU CUGAUGAG X CGAA ICUGGGUG	
1762	ACCCAGCC A AGCGGCU		CAGCCGCU CUGAUGAG X CGAA ICGUGGU	
1769	CAAGCGGC U GGGCUGUG		CACAGCCC CUGAUGAG X CGAA ICCGCUUG	
1774	GGCUGGGC U GUGGGCCU		AGGCCAC CUGAUGAG X CGAA ICCCAGCC	
1781	CUGUGGGC C UGAGGGGG		CCCCUCA CUGAUGAG X CGAA ICCCACAG	
1782	UGUGGGCC U GAGGGGGA		UCCCCUC CUGAUGAG X CGAA IGCCCACA	
1808	GAGAGAGC A UGCCUUCU		AGAAGGCA CUGAUGAG X CGAA ICUCUCUC	

Table 63

1812	GAGCAUGC C UUCUCCG		CGGAAGAA CUGAUGAG X CGAA ICAUGCUC	
1813	AGCAUGCC U UCUUCCG		CCGGAAGA CUGAUGAG X CGAA IGCAUGCU	
1816	AUGCCUUC U UCCGGAGG		CCUCCGGA CUGAUGAG X CGAA IAAGGCAU	
1819	CCUUCUUC C GGAGGAUC		GAUCCUCC CUGAUGAG X CGAA IAAGAAGG	
1831	GGAUCGAC U GGGAAAAA		UUUUUCCC CUGAUGAG X CGAA IUCGAUCC	
1841	GGAAAAAC U GGAGAACA		UGUUCUCC CUGAUGAG X CGAA IUUUUCC	
1849	UGGAGAAC A GGGAGAUC		GAUCUCCC CUGAUGAG X CGAA IUUCUCCA	
1858	GGGAGAUC C AGCCACCA		UGGUGGCU CUGAUGAG X CGAA IAUCUCCC	
1859	GGAGAUC C GCCACCAU		AUGGUGGC CUGAUGAG X CGAA IGAUCUCC	
1862	GAUCCAGC C ACCAUUCA		UGAAUGGU CUGAUGAG X CGAA ICUGGAUC	
1863	AUCCAGCC A CCAUCAA		UUGAAUGG CUGAUGAG X CGAA IGCUGGAU	
1865	CCAGCCAC C AUUCAAGC		GCUUGAAU CUGAUGAG X CGAA IUGGCUGG	
1866	CAGCCACC A UUCAAGCC		GGCUUGAA CUGAUGAG X CGAA IGUGGCUG	
1870	CACCAUUC A AGCCCAA		UUUGGGCU CUGAUGAG X CGAA IAAUGGUG	
1874	AUUCAAGC C CAAAGUGU		ACACUUUG CUGAUGAG X CGAA ICUUGAAU	
1875	UUCAAGCC C AAAGUGUG		CACACUUU CUGAUGAG X CGAA IGCUGAA	
1876	UCAAGCCC A AAGUGUGU		ACACACUU CUGAUGAG X CGAA IGGCUUGA	
1888	UGUGUGGC A AAGGAGCA		UGCUCUUU CUGAUGAG X CGAA ICCACACA	
1896	AAAGGAGC A GAGAACUU		AAGUUCUC CUGAUGAG X CGAA ICUCUUUU	
1903	CAGAGAAC U UUGACAAG		CUUGUCAA CUGAUGAG X CGAA IUUCUCUG	
1909	ACUUUGAC A AGUUCUUC		GAAGAACU CUGAUGAG X CGAA IUCAAAGU	
1915	ACAAGUUC U UCACACGA		UCGUGUGA CUGAUGAG X CGAA IAACUUGU	
1918	AGUUCUUC A CACGAGGA		UCCUCGUG CUGAUGAG X CGAA IAAGAACU	
1920	UUCUUCAC A CGAGGACA		UGUCCUCG CUGAUGAG X CGAA IUGAAGAA	
1928	ACGAGGAC A GCCCGUCU		AGACGGGC CUGAUGAG X CGAA IUCCUCGU	
1931	AGGACAGC C CGUCUUA		UUAAGACG CUGAUGAG X CGAA ICUGUCCU	
1932	GGACAGCC C GUCUUAAC		GUUAAGAC CUGAUGAG X CGAA IGCUGUCC	
1936	AGCCCGUC U UAACACCA		UGGUGUUA CUGAUGAG X CGAA IACGGGCU	
1941	GUCUUAAC A CCACCUGA		UCAGGUGG CUGAUGAG X CGAA IUUAAGAC	
1943	CUUAACAC C ACCUGAUC		GAUCAGGU CUGAUGAG X CGAA IUGUUAAG	
1944	UUAACACC A CCUGAUC		UGAUCAGG CUGAUGAG X CGAA IGUGUUA	
1946	AACACCAC C UGAUCAGC		GCUGAUC CUGAUGAG X CGAA IUGGUGUU	
1947	ACACCACC U GAUCAGCU		AGCUGAUC CUGAUGAG X CGAA IGUGGUGU	
1952	ACCUGAUC A GCUGGUUA		UAACCAGC CUGAUGAG X CGAA IAUCAGGU	
1955	UGAUCAGC U GGUUAUUG		CAAUAACC CUGAUGAG X CGAA ICUGAUC	
1965	GUUAUUGC U AACAUAGA		UCUAUGUU CUGAUGAG X CGAA ICAUAAC	
1969	UUGCUAAC A UAGACCAG		CUGGUCUA CUGAUGAG X CGAA IUUAGCAA	
1975	ACAUAGAC C AGUCUGAU		AUCAGACU CUGAUGAG X CGAA IUUAUGU	
1976	CAUAGACC A GUCUGAUU		AAUCAGAC CUGAUGAG X CGAA IGUCUAUG	
1980	GACCAGUC U GAUUUGA		UCAAAUUC CUGAUGAG X CGAA IACUGGUC	
1996	AAGGGUUC U CGUAUGUC		GACAUACG CUGAUGAG X CGAA IAACCCUU	
2005	CGUAUGUC A ACCCCAG		CUGGGGCU CUGAUGAG X CGAA IACUAACG	
2008	AUGUCAAC C CCCAGUUU		AAACUGGG CUGAUGAG X CGAA IUUGACAU	
2009	UGUCAACC C CCAGUUUG		CAAACUGG CUGAUGAG X CGAA IGUUGACA	
2010	GUCAACCC C CAGUUUGU		ACAAACUG CUGAUGAG X CGAA IGGUUGAC	
2011	UCAACCCC C AGUUUGUG		CACAAACU CUGAUGAG X CGAA IGGGUUGA	
2012	CAACCCCC A GUUUGUGC		GCACAAAC CUGAUGAG X CGAA IGGGUUG	
2021	GUUUGUGC A CCCCAUCU		AGAUGGGG CUGAUGAG X CGAA ICACAAAC	
2023	UUGUGCAC C CCAUCUUA		UAAGAUGG CUGAUGAG X CGAA IUGCACAA	
2024	UGUGCACC C CAUCUUAC		GUAAGAUG CUGAUGAG X CGAA IGUGCACA	
2025	GUGCACCC C AUCUUACA		UGUAAGAU CUGAUGAG X CGAA IGGUGCAC	

Table 63

2026	UGCACCCC A UCUIACAG		CUGUAAGA CUGAUGAG X CGAA IGGGUGCA	
2029	ACCCCAUC U UACAGAGU		ACUCUGUA CUGAUGAG X CGAA IAUGGGGU	
2033	CAUCUAC A GAGUGCAG		CUGCACUC CUGAUGAG X CGAA IUAAGAUG	
2040	CAGAGUGC A GUAUGAAA		UUUCAUAC CUGAUGAG X CGAA ICACUCUG	
2050	UAUGAAAC U CACCAGCG		CGCUGGUG CUGAUGAG X CGAA IUUUCUA	
2052	UGAAACUC A CCAGCGAG		CUCGUGG CUGAUGAG X CGAA IAGUUUCA	
2054	AAACUCAC C AGCGAGAA		UUCUCGCU CUGAUGAG X CGAA IUGAGUUU	
2055	AACUCACC A GCGAGAAC		GUUCUCGC CUGAUGAG X CGAA IGUGAGUU	
2064	GCGAGAAC A AACACCUC		GAGGUGUU CUGAUGAG X CGAA IUUCUCGC	
2068	GAACAAAC A CCUCCCCA		UGGGGAGG CUGAUGAG X CGAA IUUUGUUC	
2070	ACAAACAC C UCCCCAGC		GCUGGGGA CUGAUGAG X CGAA IUGUUUGU	
2071	CAAACACC U CCCCAGCC		GGCUGGGG CUGAUGAG X CGAA IGUGUUUG	
2073	AACACCUC C CCAGCCCC		GGGGCUGG CUGAUGAG X CGAA IAGGUGUU	
2074	ACACCUC C CAGCCCCC		GGGGGCU CUGAUGAG X CGAA IGAGGUGU	
2075	CACCUC C AGCCCCCA		UGGGGGCU CUGAUGAG X CGAA IGGAGGUG	
2076	ACUCCCC A GCCCCCAG		CUGGGGGC CUGAUGAG X CGAA IGGGAGGU	
2079	UCCCCAG C CCCAGCCC		GGGCUGGG CUGAUGAG X CGAA ICUGGGGA	
2080	CCCCAGCC C CCAGCCCU		AGGGCUGG CUGAUGAG X CGAA ICGUGGGG	
2081	CCCAGCCC C CAGCCUC		GAGGGCUG CUGAUGAG X CGAA IGGCUGGG	
2082	CCAGCCCC C AGCCUCC		GGAGGGCU CUGAUGAG X CGAA IGGGCUGG	
2083	CAGCCCCC A GCCUCCC		GGGAGGGC CUGAUGAG X CGAA IGGGGCUG	
2086	CCCCCAGC C CUCCCCGC		GCGGGGAG CUGAUGAG X CGAA ICUGGGGG	
2087	CCCCAGCC C UCCCCGCA		UGC GGGA CUGAUGAG X CGAA ICGUGGGG	
2088	CCCAGCCC U CCCCCGAG		CUGCGGGG CUGAUGAG X CGAA IGGCUGGG	
2090	CAGCCUC C CCGCAGUG		CACUGCGG CUGAUGAG X CGAA IAGGGCUG	
2091	AGCCUCC C CGCAGUGG		CCACUGCG CUGAUGAG X CGAA IGAGGGCU	
2092	GCCUCC C GCAGUGGA		UCCACUGC CUGAUGAG X CGAA IGGAGGGC	
2095	CUCCCCGC A GUGGAAGU		ACUCCAC CUGAUGAG X CGAA ICGGGGAG	
2109	AGUGAAUC C UUAACCCU		AGGGUUA CUGAUGAG X CGAA IAUUCACU	
2110	GUGAAUCC U UAACCUA		UAGGGUUA CUGAUGAG X CGAA IGAUUCAC	
2115	UCCUUAAC C CUAAAUU		AAUUUUA CUGAUGAG X CGAA IUUAAGGA	
2116	CCUUAACC C UAAAUUU		AAAUUUUA CUGAUGAG X CGAA IGUUAAGG	
2117	CUUAACCC U AAAUUUU		AAAAUUUU CUGAUGAG X CGAA IGGUUAAG	
2131	UUUAAGGC C ACGGCUUG		CAAGCCGU CUGAUGAG X CGAA ICCUUAAA	
2132	UUAAGGCC A CGGCUUGU		ACAAGCCG CUGAUGAG X CGAA IGCCUUA	
2137	GCCACGGC U UGUGUCUG		CAGACACA CUGAUGAG X CGAA ICCGUGGC	
2144	CUUGUGUC U GAUCCAU		AUGGAAUC CUGAUGAG X CGAA IACACAAG	
2150	UCUGAUUC C AUAUGGAG		CUCCAUAU CUGAUGAG X CGAA IAAUCAGA	
2151	CUGAUUCC A UAUGGAGG		CCUCCAUA CUGAUGAG X CGAA IGAAUCAG	
2161	AUGGAGGC C UGAAAAUU		AAUUUUA CUGAUGAG X CGAA ICCUCCAU	
2162	UGGAGGCC U GAAAAUUG		CAAUUUUC CUGAUGAG X CGAA IGCCUCCA	
2185	UAUAGUC C AAUUGUGA		UCACAUU CUGAUGAG X CGAA IACUAAUA	
2186	AUUAUCC A AAUGUGAU		AUCACAU CUGAUGAG X CGAA IGACUAAU	
2196	AUGUGAUC A ACUGUUA		UGAACAGU CUGAUGAG X CGAA IAUACAU	
2199	UGAUAAC U GUUCAGGG		CCCUGAAC CUGAUGAG X CGAA IUUGAUCA	
2204	AACUGUUC A GGUUCUCU		AGAGACC CUGAUGAG X CGAA IAACAGUU	
2210	UCAGGGUC U CUCUUA		UAAGAGAG CUGAUGAG X CGAA IACCCUGA	
2212	AGGGUCUC U CUUACA		UGUAAGAG CUGAUGAG X CGAA IAGACCCU	
2214	GGUCUCUC U CUUACAAC		GUUGUAAG CUGAUGAG X CGAA IAGAGACC	
2216	UCUCUCUC U UACAACCA		UGGUUGUA CUGAUGAG X CGAA IAGAGAGA	
2220	UCUCUAC A ACCAAGAA		UUCUUGGU CUGAUGAG X CGAA IUAAGAGA	

Table 63

2223	CUUACAAC C AAGAACAU		AUGUUCUU CUGAUGAG X CGAA IUUGUAAG	
2224	UUACAACC A AGAACAUU		AAUGUUCU CUGAUGAG X CGAA IGUUGUAA	
2230	CCAAGAAC A UUAUCUUA		UAAGAUAA CUGAUGAG X CGAA IUUCUUGG	
2236	ACAUUAUC U UAGUGGAA		UCCACUA CUGAUGAG X CGAA IAUAAUGU	

Input Sequence = PRKCA. Cut Site = CH/.

Stem Length = 8 . Core Sequence = CUGAUGAG X CGAA (X = GCCGUUAGGC or other stem II)

PRKCA (Homo sapiens protein kinase C, alpha (PRKCA) mRNA.; 2245 bp)

Table 64

Table 64: Activity of ribozyme core substituted analogues**A**

RYH/	All ribo I-15.1 K_{obs} (min^{-1})	G-5, A-6, G-8, G-12, I-15.1 ribo 2'-O-allyl environment K_{obs} (min^{-1})		
		U-4=ribo U	U-4=2'-amino U	U-4=2'-O-alkyl U
GCA	0.39	0.10	0.08	0.02
GCC	0.19	0.03	0.01	0.003
GCU	0.028	0.025	0.013	0.002

B

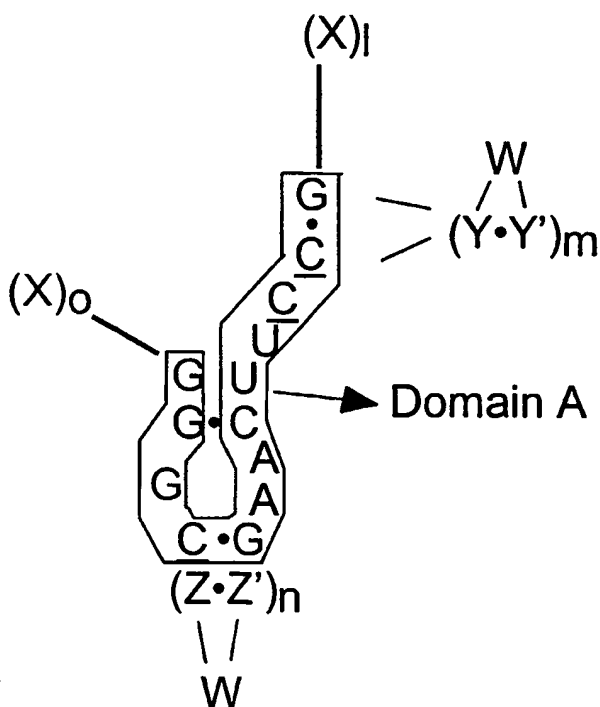
RYH/	All ribo A-15.1 K_{obs} (min^{-1})	G-5, A-6, G-8, G-12, A-15.1 ribo 2'-O-allyl environment K_{obs} (min^{-1})		
		U-4=ribo U	U-4=2'-amino U	U-4=2'-O-alkyl U
GUA	0.12	0.06	0.04	0.01
GUC	0.15	0.015	0.014	0.001
GUU	0.04	0.031	0.012	0.008

Comparison of single turnover cleavage rates for GCH and GUH substrates with I-15.1 and A-15.1 ribozymes and ribozyme analogs. Conditions: Single turnover (250 nM substrate, 2.5 μM ribozyme) pH 6.0, 37 °C, 10 mM Mg^{++}

Claims:

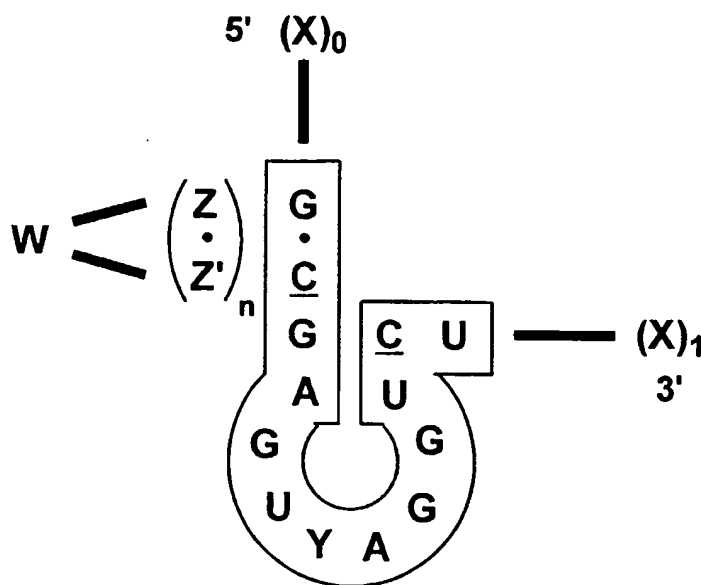
We claim:

- 5 1. An enzymatic nucleic acid molecule having formula 4 namely:



wherein each X, Y, and Z represents independently a nucleotide which may be the same or different; l is an integer greater than or equal to 3; m is an integer greater than 10 1; n is an integer greater than 1; O is an integer greater than or equal to 3; Z' is a nucleotide complementary to Z; Y' is a nucleotide complementary to Y; each X(l) and X(o) are oligonucleotides which are of sufficient length to stably interact independently with a target nucleic acid sequence; W is a linker of ≥ 2 nucleotides; A, U, G, and C represent nucleotides; C is 2'-amino; and ____ represents a chemical 15 linkage.

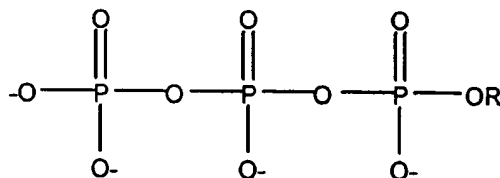
2. An enzymatic nucleic acid molecule having formula 5 namely:



- 5 wherein each X, Y, and Z represents independently a nucleotide which may be the same or different; l is an integer greater than or equal to 3; n is an integer greater than 1; 0 is an integer greater than or equal to 3; Z' is a nucleotide complementary to Z; each X_(l) and X_(o) are oligonucleotides which are of sufficient length to stably interact independently with a target nucleic acid sequence; W is a linker of ≥ 2 nucleotides in
 10 length or may be a non-nucleotide linker; A, U, G, and C represent nucleotides; C is 2'-amino; and ___ represents a chemical linkage.
3. The enzymatic nucleic acid molecule of claims 1 or 2, wherein l is selected from the group consisting of 4, 5, 6, 7, 8, 9, 10, 11, 12, and 15.
4. The enzymatic nucleic acid molecule of claim 1, wherein m is selected from the
 15 group consisting of 2, 3, 4, 5, 6, and 7.
5. The enzymatic nucleic acid molecule of claims 1 or 2, wherein n is selected from the group consisting of 2, 3, 4, 5, 6, and 7.
6. The enzymatic nucleic acid molecule of claims 1 or 2, wherein o is selected from the group consisting of 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, and 15.

7. The enzymatic nucleic acid molecule of claims 1 or 2, wherein l and o are of the same length.
8. The enzymatic nucleic acid molecule of claims 1 or 2, wherein l and o are of different length.
- 5 9. The enzymatic nucleic acid molecule of claims 1 or 2, wherein the target nucleic acid sequence is selected from the group consisting of an RNA, DNA and RNA/DNA mixed polymer.
10. The enzymatic nucleic acid molecule of claims 1 or 2, wherein said chemical linkage is selected from the group consisting of phosphate ester linkage, amide
10 linkage, phosphorothioate, and phosphorodithioate.
11. The enzymatic nucleic acid molecule of claims 1 or 2, wherein said C_{12} is selected from the group consisting of 2'-deoxy-2'-NH₂ and 2'-deoxy-2'-O-NH₂.
12. A method for inhibiting expression of a gene in a cell, comprising the step of administering to said cell the enzymatic nucleic acid molecule of claims 1 or 2
15 under conditions suitable for said inhibition.
13. A method of cleaving a separate RNA molecule comprising, contacting the enzymatic nucleic acid molecule of claims 1 or 2 with said separate RNA molecule under conditions suitable for the cleavage of said separate RNA molecule.
14. The method of claim 13, wherein said cleavage is carried out in the presence of a
20 divalent cation.
15. The method of claim 14, wherein said divalent cation is Mg²⁺.
16. The enzymatic nucleic acid molecule of claims 1 or 2, wherein said enzymatic nucleic acid molecule is chemically synthesized.
17. The enzymatic nucleic acid molecule of claims 1 or 2, wherein said enzymatic
25 nucleic acid molecule comprises at least one ribonucleotide.
18. The enzymatic nucleic acid molecule of claims 1 or 2, wherein said enzymatic nucleic acid molecule comprises no ribonucleotide residues.
19. The enzymatic nucleic acid molecule of claims 1 or 2, wherein said enzymatic nucleic acid molecule comprises at least one 2-amino modification.

20. The enzymatic nucleic acid molecule of claims 1 or 2, wherein said enzymatic nucleic acid molecule comprises at least three phosphorothioate modifications.
21. The enzymatic nucleic acid molecule of claim 20, wherein said phosphorothioate modification is at the 5'-end of said enzymatic nucleic acid molecule.
- 5 22. The enzymatic nucleic acid molecule of claims 1 or 2, wherein said enzymatic nucleic acid molecule comprises a 5'-cap or a 3'-cap or both a 5'-cap and a 3'-cap.
23. The enzymatic nucleic acid molecule of claim 22, wherein said 5-cap is phosphorothioate modification.
24. The enzymatic nucleic acid molecule of claim 22, wherein said 3'-cap is an
10 inverted abasic moiety.
25. A compound having the formula 3:



- wherein R is independently any nucleoside selected from the group consisting of 2'-O-methyl-2,6-diaminopurine riboside; 2'-deoxy-2'-amino-2,6-diaminopurine riboside; 2'-
15 (N-alanyl) amino-2'-deoxy-uridine; 2'-(N-phenylalanyl)amino-2'-deoxy-uridine; 2'-deoxy-2'-(N-β-alanyl) amino ; 2'-deoxy-2'-(lysiyl) amino uridine; 2'-C-allyl uridine; 2'-O-amino-uridine; 2'-O-methylthiomethyl adenosine; 2'-O-methylthiomethyl cytidine ; 2'-O-methylthiomethyl guanosine; 2'-O-methylthiomethyl-uridine; 2'-deoxy-2'-(N-histidyl) amino uridine; 2'-deoxy-2'-amino-5-methyl cytidine; 2'-(N-β-carboxamidine-β-alanyl)amino-2'-deoxy-uridine; 2'-deoxy-2'-(N-β-alanyl)-guanosine; 2'-O-amino-adenosine; 2'-(N-lysyl)amino-2'-deoxy-cytidine; 2'-Deoxy-2'-(L-histidine) amino Cytidine; 5-Imidazoleacetic acid 2'-deoxy uridine, 5-[3-(N-4-imidazoleacetyl)aminopropynyl]-2'-O-methyl uridine, 5-(3-aminopropynyl)-2'-O-methyl uridine, 5-(3-aminopropyl)-2'-O-methyl uridine, 5-[3-(N-4-imidazoleacetyl)aminopropyl]-2'-O-methyl uridine, 5-(3-aminopropyl)-2'-deoxy-2'-fluoro uridine, 2'-Deoxy-2'-(β-alanyl-L-histidyl)amino uridine, 2'-deoxy-2'-β-alaninamido-uridine, 3-(2'-deoxy-2'-fluoro-β-D-ribofuranosyl)piperazino[2,3-D]pyrimidine-2-one, 5-[3-(N-4-imidazoleacetyl)aminopropyl]-2'-deoxy-2'-fluoro uridine, 5-[3-(N-4-imidazoleacetyl)aminopropynyl]-2'-deoxy-2'-fluoro uridine, 5-E-
- 20
- 25

(2-carboxyvinyl-2'-deoxy-2'-fluoro uridine, 5-[3-(N-4-aspartyl)aminopropynyl-2'-fluoro uridine, 5-(3-aminopropyl)-2'-deoxy-2-fluoro cytidine, and 5-[3-(N-4-succinyl)aminopropyl-2'-deoxy-2-fluoro cytidine.

26. A process for incorporation of the compounds of claim 25 into an oligonucleotide
5 comprising the step of contacting said compound with a mixture comprising a nucleic acid template, an RNA polymerase enzyme, and an enhancer of modified nucleotide triphosphate incorporation, under conditions suitable for the incorporation of said compound into said oligonucleotide.
27. The process of claim 26, wherein said RNA polymerase is a T7 RNA polymerase.
- 10 28. The process of claim 26, wherein said RNA polymerase is a mutant T7 RNA polymerase.
29. The process of claim 26, wherein said RNA polymerase is a SP6 RNA polymerase.
30. The process of claim 26, wherein said RNA polymerase is a mutant SP6 RNA polymerase.
- 15 31. The process of claim 26, wherein said RNA polymerase is a T3 RNA polymerase.
32. The process of claim 26, wherein said RNA polymerase is a mutant T3 RNA polymerase.
33. The process of claim 26, wherein said enhancer of modified nucleotide triphosphate incorporation is selected from the group consisting of LiCl, methanol, polyethylene glycol, diethyl ether, propanol, methylamine, and ethanol.
20
34. A process for the synthesis of a pyrimidine nucleotide triphosphate comprising the steps of:
- a. monophosphorylation, wherein a pyrimidine nucleoside is contacted with a mixture comprising a phosphorylating reagent, a trialkyl phosphate and dimethylaminopyridine, under conditions suitable for the formation of a pyrimidine
25 nucleotide monophosphate; and
- b. pyrophosphorylation, wherein said pyrimidine monophosphate from step (a) is contacted with a pyrophosphorylating reagent under conditions suitable for the formation of said pyrimidine nucleotide triphosphate.

35. The process of claim 34, wherein said pyrimidine nucleoside triphosphate is uridine triphosphate.
36. The process of claim 34, wherein said uridine triphosphate has a 2'-sugar modification.
- 5 37. The process of claim 36, wherein said uridine triphosphate is 2'-O-methylthiomethyl uridine triphosphate.
38. The process of claim 34, wherein said phosphorylating agent is selected from the group consisting of phosphorus oxychloride, phospho-tris-triazolides and phospho-tris-triimidazolides.
- 10 39. The process of claim 34, wherein said trialkylphosphate is triethyl phosphate.
40. The process of claim 34, wherein said pyrophosphorylating reagent is tributyl ammonium pyrophosphate.
41. The process of claim 26, wherein said oligonucleotide is RNA.
42. The process of claim 26, wherein said oligonucleotide is an enzymatic nucleic acid
15 molecule.
43. The process of claim 26, wherein said oligonucleotide is an aptamer.
44. A kit for synthesis of an oligonucleotide comprising an RNA polymerase, an enhancer of modified nucleotide triphosphate incorporation and at least one compound of claim 25.
- 20 45. A kit for synthesis of an oligonucleotide comprising a DNA polymerase, an enhancer of modified nucleotide triphosphate incorporation and at least one compound of claim 25.
46. The kit of claim 44, wherein said RNA polymerase is a bacteriophage T7 RNA polymerase.
- 25 47. The kit of claim 44, wherein said RNA polymerase is a bacteriophage SP6 RNA polymerase.
48. The kit of claim 44, wherein said RNA polymerase is a bacteriophage T3 RNA polymerase.

49. The kit of claim 44, wherein said RNA polymerase is a mutant T7 RNA polymerase.
50. The kit of claim 44 or 45, wherein said kit comprises at least two different compounds of claim 25.
- 5 51. A nucleic acid catalyst comprising a histidyl modification, wherein said nucleic acid catalyst is able to catalyze an endonuclease reaction in the absence of a metal ion co-factor.
52. The nucleic acid catalyst of claim 51, wherein said catalyst is able to cleave a separate nucleic acid molecule.
- 10 53. The nucleic acid catalyst of claim 52, wherein said separate nucleic acid molecule is an RNA molecule.
54. The nucleic acid catalyst of claim 52, wherein said separate nucleic acid molecule is a DNA molecule.
- 15 55. The nucleic acid catalyst of claim 51, wherein said nucleic acid catalyst comprises at least one ribonucleotide.
56. The enzymatic nucleic acid molecule of claim 2, wherein said nucleic acid molecule has an endonuclease activity to cleave RNA of HER2 gene.
57. The enzymatic nucleic acid molecule of claim 56, wherein said nucleic acid molecule comprises sequences complementary to any of substrate sequences
20 defined as Target sequence in Tables 58, 59 and 62.
58. The enzymatic nucleic acid molecule of claim 56, wherein said nucleic acid molecule comprises any of ribozyme sequences defined as Ribozyme sequence in Tables 58, 59 and 62.
- 25 59. A method for treating cancer using the enzymatic nucleic acid molecule of claim 56.
60. The method of claim 59, wherein said cancer is breast cancer.
61. A method for treating conditions associated with the level of HER2 gene using the enzymatic nucleic acid molecule of claim 56.

62. The enzymatic nucleic acid molecule of claim 56, wherein said enzymatic nucleic acid molecule comprises a substrate binding region which has between 5 and 30 nucleotides complementary to the RNA.
- 5 63. The enzymatic nucleic acid molecule of claim 56, wherein said enzymatic nucleic acid molecule comprises a substrate binding region which has between 7 and 12 nucleotides complementary to the RNA.
64. A mammalian cell including the enzymatic nucleic acid molecule of claim 56.
65. The mammalian cell of claim 64, wherein said mammalian cell is a human cell.
66. A mammalian cell including the enzymatic nucleic acid molecule of claims 1 or 2.
- 10 67. The mammalian cell of claim 66, wherein said mammalian cell is a human cell.
68. A method for inhibiting expression of HER2 gene in a cell, comprising the step of administering to said cell the enzymatic nucleic acid molecule of claim 56 under conditions suitable for said inhibition.
- 15 69. A method of cleaving RNA derived from HER2 gene comprising, contacting the enzymatic nucleic acid molecule of claim 56 with said RNA molecule under conditions suitable for the cleavage of said RNA molecule.
70. A pharmaceutical composition comprising the enzymatic nucleic acid molecule of any of claims 1 or 2.
- 20 71. A pharmaceutical composition comprising the enzymatic nucleic acid molecule of claim 56.
72. A method of treatment of a patient having a condition associated with the level of HER2, wherein said patient is administered the enzymatic nucleic acid molecule of claim 56 under conditions suitable for said treatment.
- 25 73. The method of claim 72, wherein said method is performed in conjunction with one or more other therapies.
74. The method of claim 59, wherein said enzymatic nucleic acid molecule is used in conjunction with one or more other therapies.
75. The enzymatic nucleic acid molecule of claim 56, wherein said enzymatic nucleic acid molecule comprises at least one sugar modification.

76. The enzymatic nucleic acid molecule of claim 56, wherein said enzymatic nucleic acid molecule comprises at least one nucleic acid base modification.
77. The enzymatic nucleic acid molecule of claim 56, wherein said enzymatic nucleic acid molecule comprises at least one phosphate backbone modification.
- 5 78. The enzymatic nucleic acid molecule of claim 56, wherein said phosphate backbone modification is selected from the group consisting of phosphorothioate, phosphorodithioate and amide.
79. An enzymatic nucleic acid molecule which down regulates expression of genes selected from the group consisting of beta site APP-cleaving enzyme (BACE) and
10 telomerase reverse transcriptase (TERT) genes.
80. The enzymatic nucleic acid molecule of claim 79, wherein said gene is the beta site APP-cleaving enzyme (BACE).
81. The enzymatic nucleic acid molecule of claim 79, wherein said gene is the telomerase reverse transcriptase (TERT).
- 15 82. A nucleic acid molecule which down regulates expression of genes selected from the group consisting of protein-tyrosine phosphatase-1B (PTP-1B), methionine aminopeptidase (MetAP-2), hepatitis B virus (HBV), phospholamban (PLN), and presenilin (ps-2) genes.
83. The nucleic acid molecule of claim 82, wherein said nucleic acid molecule is an
20 enzymatic nucleic acid molecule.
84. The nucleic acid molecule of claim 82, wherein said nucleic acid molecule is an antisense nucleic acid molecule.
85. The nucleic acid molecule of any of claims 82-84, wherein said gene is the protein-tyrosine phosphatase-1B (PTP-1B).
- 25 86. The nucleic acid molecule of any of claims 82-84, wherein said gene is the methionine aminopeptidase (MetAP-2).
87. The nucleic acid molecule of any of claims 82-84, wherein said gene is the hepatitis B virus (HBV).

88. The nucleic acid molecule of any of claims 82-84, wherein said gene is the phospholamban (PLN).
89. The nucleic acid molecule of any of claims 82-84, wherein said gene is the presenilin (ps-2).
- 5 90. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid molecule is adapted for use to treat diseases and conditions related to the expression of genes selected from the group consisting of beta site APP-cleaving enzyme (BACE), telomerase reverse transcriptase (TERT), protein-tyrosine phosphatase-1B (PTP-1B), methionine aminopeptidase (MetAP-2),
10 hepatitis B virus (HBV), phospholamban (PLN), and presenilin (ps-2) genes.
91. The nucleic acid molecule of claim 82, wherein said nucleic acid molecule is adapted for use to treat diseases and conditions related to the expression of genes selected from the group consisting of protein-tyrosine phosphatase-1B (PTP-1B), methionine aminopeptidase (MetAP-2), hepatitis B virus (HBV), phospholamban
15 (PLN), and presenilin (ps-2) genes.
92. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid molecule has an endonuclease activity to cleave RNA encoded by said beta site APP-cleaving enzyme (BACE), telomerase reverse transcriptase (TERT), protein-tyrosine phosphatase-1B (PTP-1B), methionine
20 aminopeptidase (MetAP-2), hepatitis B virus (HBV), phospholamban (PLN), and presenilin (ps-2) genes.
93. The enzymatic nucleic acid of any of claims 79 or 83, wherein a binding arm of said enzymatic nucleic acid molecule comprise sequences complementary to any of the sequences defined as Target or Substrate sequence in Tables 3-30, and 36-
25 43.
94. The enzymatic nucleic acid molecule of any of claims 79 or 83 wherein said enzymatic nucleic acid molecule comprises any of the sequences defined as Ribozyme or DNAzyme sequence in Tables 3-29, and 37-43.
95. The nucleic acid molecule of claim 84, wherein said antisense nucleic acid
30 molecule comprises sequence complementary to any of the sequences defined as Target or Substrate sequence in Tables 3-12, 24-30, and 36-43.

96. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid molecule is in a hammerhead (HH) motif.
97. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid molecule is in a zinzyme (Class II) motif.
- 5 98. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid molecule is in an amberzyme (Class I) motif.
99. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid molecule is in a hairpin, hepatitis Delta virus, group I intron, VS nucleic acid, or RNase P nucleic acid motif.
- 10 100. The enzymatic nucleic acid molecule of claim 97, wherein said zinzyme motif comprises sequences complementary to any of the substrate sequences shown in Tables 21, 27 and 40.
101. The enzymatic nucleic acid molecule of claim 98, wherein said amberzyme motif comprises sequences complementary to any of the substrate sequences
15 shown in Tables 23, 29, and 42.
102. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid molecule is in a NCH motif.
103. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid molecule is in a G-cleaver motif.
- 20 104. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid molecule is a DNAzyme.
105. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid molecule comprises between 12 and 100 bases complementary to the RNA of genes selected from the group consisting of beta site
25 APP-cleaving enzyme (BACE), telomerase reverse transcriptase (TERT), protein-tyrosine phosphatase-1B (PTP-1B), methionine aminopeptidase (MetAP-2), hepatitis B virus (HBV), phospholamban (PLN), and presenilin (ps-2) genes.
106. The enzymatic nucleic acid of any of claims 79 or 83, wherein said enzymatic nucleic acid molecule comprises between 14 and 24 bases complementary to the
30 RNA of genes selected from the group consisting of beta site APP-cleaving enzyme (BACE), telomerase reverse transcriptase (TERT), protein-tyrosine

phosphatase-1B (PTP-1B), methionine aminopeptidase (MetAP-2), hepatitis B virus (HBV), phospholamban (PLN), and presenilin (ps-2) genes.

107. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid is chemically synthesized.
- 5 108. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid comprises at least one 2'-sugar modification.
109. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid comprises at least one nucleic acid base modification.
- 10 110. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid comprises at least one phosphate backbone modification.
111. A mammalian cell including the enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said mammalian cell is not a living human.
112. The mammalian cell of claim 111, wherein said mammalian cell is a human cell.
- 15 113. The antisense nucleic acid molecule of claim 84, wherein said antisense nucleic acid is chemically synthesized.
114. The antisense nucleic acid molecule of claim 84, wherein said antisense nucleic acid comprises at least one 2'-sugar modification.
- 20 115. The antisense nucleic acid molecule of claim 84, wherein said antisense nucleic acid comprises at least one nucleic acid base modification.
116. The antisense nucleic acid molecule of claim 84, wherein said antisense nucleic acid comprises at least one phosphate backbone modification.
117. A mammalian cell including the antisense nucleic acid molecule of claim 84, wherein said mammalian cell is not a living human.
- 25 118. The mammalian cell of claim 117, wherein said mammalian cell is a human cell.
119. A method of reducing BACE activity in a cell, comprising the step of contacting said cell with the enzymatic nucleic acid molecule of claim 80, under conditions suitable for said inhibition.

120. A method of reducing TERT activity in a cell, comprising the step of contacting said cell with the enzymatic nucleic acid molecule of claim 81, under conditions suitable for said inhibition.
- 5 121. A method of reducing PTP-1B activity in a cell, comprising the step of contacting said cell with the nucleic acid molecule of claim 85, under conditions suitable for said inhibition.
122. A method of reducing MetAP-2 activity in a cell, comprising the step of contacting said cell with the nucleic acid molecule of claim 86, under conditions suitable for said inhibition.
- 10 123. A method of reducing HBV activity in a cell, comprising the step of contacting said cell with the nucleic acid molecule of claim 87, under conditions suitable for said inhibition.
124. A method of reducing phospholamban (PLN) activity in a cell, comprising the step of contacting said cell with the nucleic acid molecule of claim 88, under
15 conditions suitable for said inhibition.
125. A method of reducing presenilin-2 (ps-2) activity in a cell, comprising the step of contacting said cell with the nucleic acid molecule of claim 89, under conditions suitable for said inhibition.
- 20 126. A method of treatment of a patient having a condition associated with the level of BACE, comprising contacting cells of said patient with the enzymatic nucleic acid molecule of claim 80, under conditions suitable for said treatment.
127. A method of treatment of a patient having a condition associated with the level of TERT, comprising contacting cells of said patient with the enzymatic nucleic acid molecule of claim 81, under conditions suitable for said treatment.
- 25 128. A method of treatment of a patient having a condition associated with the level of PTP-1B, comprising contacting cells of said patient with the nucleic acid molecule of claim 85, under conditions suitable for said treatment.
- 30 129. A method of treatment of a patient having a condition associated with the level of MetAP-2, comprising contacting cells of said patient with the nucleic acid molecule of claim 86, under conditions suitable for said treatment.

130. A method of treatment of a patient having a condition associated with the level of HBV, comprising contacting cells of said patient with the nucleic acid molecule of claim 87, under conditions suitable for said treatment.
131. A method of treatment of a patient having a condition associated with the level
5 of phospholamban (PLN), comprising contacting cells of said patient with the nucleic acid molecule of claim 88, under conditions suitable for said treatment.
132. A method of treatment of a patient having a condition associated with the level of presenilin-2 (ps-2), comprising contacting cells of said patient with the nucleic acid molecule of claim 89, under conditions suitable for said treatment.
- 10 133. The method of any of claims 126-132 further comprising the use of one or more drug therapies under conditions suitable for said treatment.
134. A method of cleaving RNA of BACE gene, comprising, contacting the enzymatic nucleic acid molecule of claim 80, with said RNA under conditions suitable for the cleavage of said RNA.
- 15 135. A method of cleaving RNA of TERT gene, comprising, contacting the enzymatic nucleic acid molecule of claim 81, with said RNA under conditions suitable for the cleavage of said RNA.
136. A method of cleaving RNA of PTP-1B gene, comprising, contacting the enzymatic nucleic acid molecule of claim 85, with said RNA under conditions
20 suitable for the cleavage of said RNA.
137. A method of cleaving RNA of MetAP-2 gene, comprising, contacting the enzymatic nucleic acid molecule of claim 86, with said RNA under conditions suitable for the cleavage of said RNA.
138. A method of cleaving RNA of HBV gene, comprising, contacting the
25 enzymatic nucleic acid molecule of claim 87, with said RNA under conditions suitable for the cleavage of said RNA.
139. A method of cleaving RNA of phospholamban (PLN) gene, comprising, contacting the enzymatic nucleic acid molecule of claim 88, with said RNA under conditions suitable for the cleavage of said RNA.

140. A method of cleaving RNA of presenilin-2 (ps-2) gene, comprising, contacting the enzymatic nucleic acid molecule of claim 89, with said RNA under conditions suitable for the cleavage of said RNA.
141. The method of any of claims 134-140, wherein said cleavage is carried out in the presence of a divalent cation.
142. The method of claim 141, wherein said divalent cation is Mg^{2+} .
143. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid comprises a cap structure, wherein the cap structure is at the 5'-end or 3'-end or both the 5'-end and the 3'-end.
144. The antisense nucleic acid molecule of claim 84, wherein said antisense nucleic acid comprises a cap structure, wherein the cap structure is at the 5'-end or 3'-end or both the 5'-end and the 3'-end.
145. The enzymatic nucleic acid molecule of claim 96, wherein said hammerhead motif comprises sequences complementary to any of sequences defined as Target or Substrate sequences in Tables 3, 9, 13, 18, 24, and 37.
146. The enzymatic nucleic acid molecule of claim 102, wherein said NCH motif comprises sequences complementary to any of sequences defined as Target or Substrate sequences in Tables 4, 10, 14, 19, 25, and 38.
147. The enzymatic nucleic acid molecule of claim 103, wherein said G-cleaver motif comprises sequences complementary to any of sequences defined as Target or Substrate sequences in Tables 5, 11, 15, 20, 26, and 39.
148. The enzymatic nucleic acid molecule of claim 104, wherein said DNAzyme comprises sequences complementary to any of sequences defined as Target or Substrate sequences in Tables 6, 16, 22, 28, and 41.
149. The method of any of claims 119-125 or 133, wherein said enzymatic nucleic acid molecule is in a hammerhead motif.
150. The method of any of claims 119-125 or 133, wherein said nucleic acid molecule is a DNAzyme.

151. An expression vector comprising nucleic acid sequence encoding at least one enzymatic nucleic acid molecule of any of claims 79 or 83, in a manner which allows expression of that enzymatic nucleic acid molecule.
152. An expression vector comprising nucleic acid sequence encoding at least one
5 antisense nucleic acid molecule of claim 84, in a manner which allows expression of that antisense nucleic acid molecule.
153. A mammalian cell including an expression vector of any of claims 151 or 152, wherein said mammalian cell is not a living human.
154. The mammalian cell of claim 153, wherein said mammalian cell is a human
10 cell.
155. The expression vector of claim 151, wherein said enzymatic nucleic acid molecule is in a hammerhead motif.
156. The expression vector of claim 151, wherein said expression vector further
15 comprises a sequence for an antisense nucleic acid molecule complementary to the RNA of genes selected from the group consisting of beta site APP-cleaving enzyme (BACE), telomerase reverse transcriptase (TERT), protein-tyrosine phosphatase-1B (PTP-1B), methionine aminopeptidase (MetAP-2), hepatitis B virus (HBV), phospholamban (PLN), and presenilin (ps-2) genes.
157. The expression vector of claim 151, wherein said expression vector comprises
20 sequence encoding at least two said enzymatic nucleic acid molecules, which may be same or different.
158. The expression vector of claim 157, wherein one said expression vector further
25 comprises sequence encoding antisense nucleic acid molecule complementary to the RNA of genes selected from the group consisting of beta site APP-cleaving enzyme (BACE), telomerase reverse transcriptase (TERT), protein-tyrosine phosphatase-1B (PTP-1B), methionine aminopeptidase (MetAP-2), hepatitis B virus (HBV), phospholamban (PLN), and presenilin (ps-2) genes.
159. A method for treatment of Alzheimer's disease comprising the step of
30 administering to a patient the enzymatic nucleic acid molecule of claim 80 under conditions suitable for said treatment.

160. The method of claim 159, wherein said treatment of Alzheimer's disease is treatment of dementia.
161. A method for treatment of Alzheimer's disease comprising the step of administering to a patient the antisense nucleic acid molecule of claim 89 under conditions suitable for said treatment.
162. A method for treatment of diabetes comprising the step of administering to a patient the nucleic acid molecule of claim 85 under conditions suitable for said treatment.
163. The method of claim 162, wherein said diabetes is type I diabetes.
164. The method of claim 162, wherein said diabetes is type II diabetes.
165. A method for treatment of diabetes comprising the step of administering to a patient the antisense nucleic acid molecule of claim 85 under conditions suitable for said treatment.
166. A method for treatment of obesity comprising the step of administering to a patient the nucleic acid molecule of claim 85 under conditions suitable for said treatment.
167. A method for treatment of obesity comprising the step of administering to a patient the antisense nucleic acid molecule of claim 85 under conditions suitable for said treatment.
168. A method for treatment of heart disease comprising the step of administering to a patient the nucleic acid molecule of claim 88 under conditions suitable for said treatment.
169. The method of claim 168, wherein said heart disease is heart failure.
170. The method of claim 168, wherein said heart disease is congestive heart failure.
171. A method for treatment of pressure overload hypertrophy, or dilated cardiomyopathy, or both, comprising the step of administering to a patient the nucleic acid molecule of claim 88 under conditions suitable for said treatment.

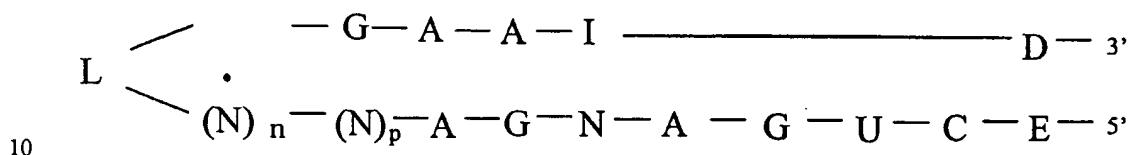
172. A method for treatment of cancer comprising the step of administering to a patient the nucleic acid molecule of claim 86 under conditions suitable for said treatment.
- 5 173. A method for treatment of hepatitis comprising the step of administering to a patient the nucleic acid molecule of claim 87 under conditions suitable for said treatment.
174. A method for treatment of hepatocellular carcinoma comprising the step of administering to a patient the nucleic acid molecule of claim 87 under conditions suitable for said treatment.
- 10 175. The method of claim 159, wherein said enzymatic nucleic acid molecule is in a hammerhead motif.
176. The method of claim 159, wherein said method further comprises administering to said patient the enzymatic nucleic acid molecule in conjunction with one or more of other therapies.
- 15 177. The method of any of claims 162, 165-168, or 171-174, wherein said nucleic acid molecule is an enzymatic nucleic acid molecule.
178. The method of any of claims 162, 166-168, or 171-174, wherein said nucleic acid molecule is an antisense nucleic acid molecule.
- 20 179. The method of any of claims 162, 165-168, or 171-174, wherein said method further comprises administering to said patient the nucleic acid molecule in conjunction with one or more of other therapies.
180. The enzymatic nucleic acid molecule of any of claims 79 or 83, wherein said enzymatic nucleic acid molecule comprises at least five ribose residues; at least ten 2'-O-methyl modifications, and a 3'- end modification.
- 25 181. The enzymatic nucleic acid molecule of claim 180, wherein said enzymatic nucleic acid molecule further comprises phosphorothioate linkages on at least three of the 5' terminal nucleotides.
182. The enzymatic nucleic acid molecule of claim 180, wherein said 3'- end modification is 3'-3' inverted abasic moiety.

183. The enzymatic nucleic acid molecule of claim 104, wherein said DNAzyme comprises at least ten 2'-O-methyl modifications and a 3'-end modification.

184. The enzymatic nucleic acid molecule of claim 183, wherein said DNAzyme further comprises phosphorothioate linkages on at least three of the 5' terminal nucleotides.

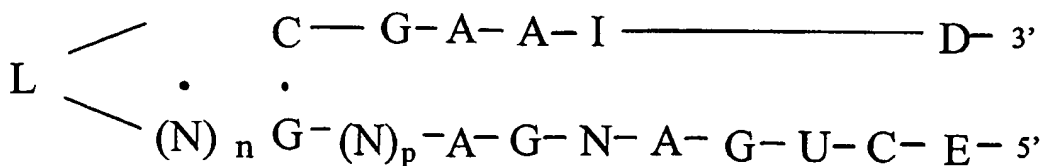
185. The enzymatic nucleic acid molecule of claim 183, wherein said 3'- end modification is 3'-3' inverted abasic moiety.

186. An enzymatic nucleic acid molecule having formula 1:



wherein N represents independently a nucleotide or a non-nucleotide linker, which may be same or different; D and E are independently oligonucleotides of length sufficient to stably interact with a target RNA molecule; o and n are integers independently greater than or equal to 1, wherein if (N)_o and (N)_n are nucleotides, (N)_o and (N)_n are optionally able to interact by hydrogen bond interaction; • indicates base-paired interaction; L is a linker which may be present or absent, but when present, is a nucleotide linker, a non-nucleotide linker, or a combination of nucleotide and a non-nucleotide linker; p is an integer 0 or 1; represents a chemical linkage; and A, U, I, C and G represent adenosine, uridine, inosine, cytidine and guanosine nucleotides, respectively.

187. An enzymatic nucleic acid molecule having formula 2:



wherein N represents independently a nucleotide or a non-nucleotide linker, which may be same or different; D and E are independently oligonucleotides of length sufficient to stably interact with a target RNA molecule; o and n are integers independently greater than or equal to 0, wherein if (N)_o and (N)_n are nucleotides,

(N)_o and (N)_n are optionally able to interact by hydrogen bond interaction; • indicates base-paired interaction; L is a linker which may be present or absent, but when present, is a nucleotide linker, a non-nucleotide linker, or a combination of nucleotide and a non-nucleotide linker; p is an integer 0 or 1; represents a chemical linkage; and A, U, I, C and G represent adenosine, uridine, inosine, cytidine and guanosine nucleotides, respectively.

188. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said D and E are independently of length selected from the group consisting of 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 17, and 20 nucleotides.
189. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said D and E are of the same length.
190. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said D and E are of different length.
191. The enzymatic nucleic acid molecule of claim 186, wherein said o and n are independently integers selected from the group consisting of 2, 3, 4, 5, 6, 7, 8, 9, 10, 15, 20, 25, 30, 35, and 50.
192. The enzymatic nucleic acid molecule of claim 187, wherein said o and n are independently integers selected from the group consisting of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 15, 20, 25, 30, 35, and 50.
193. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said (N)_o and (N)_n comprise nucleotides that are complementary to each other.
194. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said (N)_o and (N)_n are of the same length.
195. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said (N)_o and (N)_n are of different length.
196. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said L is a nucleotide linker.
197. The enzymatic nucleic acid molecule of claim 196, wherein said nucleotide linker is of length between 3-50 nucleotides.

198. The enzymatic nucleic acid molecule of claim 196, wherein said nucleotide linker is an aptamer.
199. The enzymatic nucleic acid molecule of claim 196 wherein said nucleotide linker is selected from the group consisting of 5'-GAAA-3' and 5'-GUUA-3'.
- 5 200. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said L is a non-nucleotide linker.
201. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said chemical linkage is independently or in combination selected from the group consisting of phosphate ester linkage, amide linkage, phosphorothioate, arabino,
10 arabinofluoro, and phosphorodithioate.
202. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said p is 1.
203. The enzymatic nucleic acid molecule of claim 202, wherein said N of (N)_p is independently selected from the group consisting of adenosine, uridine, and cytidine.
- 15 204. The enzymatic nucleic acid molecule of claims 186 or 187 wherein said enzymatic nucleic acid molecule is chemically synthesized.
205. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said enzymatic nucleic acid molecule comprises at least three ribonucleotide residues.
206. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said
20 enzymatic nucleic acid molecule comprises at least four ribonucleotide residues.
207. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said enzymatic nucleic acid molecule comprises at least five ribonucleotide residues.
208. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said I is selected from the group consisting of ribo-inosine and xylo-inosine.
- 25 209. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said enzymatic nucleic acid molecule comprises at least one sugar modification.
210. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said enzymatic nucleic acid molecule comprises at least nucleic acid base modification.

211. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said enzymatic nucleic acid molecule comprises at least one phosphate backbone modification.
212. The enzymatic nucleic acid molecule of claim 209, wherein said sugar
5 modification is selected from the group consisting of 2'-H, 2'-O-methyl, 2'-O-allyl, and 2'-deoxy-2'-amino.
213. The enzymatic nucleic acid molecule of claim 211, wherein said phosphate backbone modification is selected from the group consisting of phosphorothioate, phosphorodithioate and amide.
- 10 214. The enzymatic nucleic acid molecule of claims 186 or 187 wherein said enzymatic nucleic acid molecule comprises a 5'-cap or a 3'-cap or both a 5'-cap and a 3'-cap.
215. The enzymatic nucleic acid molecule of claim 214, wherein said 5'-cap is a
15 phosphorothioate modification of at least one 5'-terminal nucleotide in said enzymatic nucleic acid molecule.
216. The enzymatic nucleic acid molecule of claim 214, wherein said 5'-cap is a phosphorothioate modification of at least two 5'-terminal nucleotide in said enzymatic nucleic acid molecule.
217. The enzymatic nucleic acid molecule of claim 214, wherein said 5'-cap is a
20 phosphorothioate modification of at least three 5'-terminal nucleotide in said enzymatic nucleic acid molecule.
218. The enzymatic nucleic acid molecule of claim 214, wherein said 3'-cap is a 3'-3' inverted abasic moiety.
219. The enzymatic nucleic acid molecule of claim 214, wherein said 3'-cap is a 3'-
25 3' inverted nucleotide moiety.
220. A method for inhibiting expression of a gene in a cell, comprising the step of administering to said cell the enzymatic nucleic acid molecule of claims 186 or 187 under conditions suitable for said inhibition.
221. A method of cleaving a separate RNA molecule comprising, contacting the
30 enzymatic nucleic acid molecule of claims 186 or 187 with said separate RNA

molecule under conditions suitable for the cleavage of said separate RNA molecule.

222. The method of claim 221, wherein said cleavage is carried out in the presence of a divalent cation.

5 223. The method of claim 222, wherein said divalent cation is Mg^{2+} .

224. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said enzymatic nucleic acid molecule has an endonuclease activity to cleave RNA derived from HER2 gene.

10 225. The enzymatic nucleic acid molecule of claim 224, wherein said enzymatic nucleic acid molecule comprises sequences complementary to any of NCH substrate sequence of Table 34.

226. The enzymatic nucleic acid molecule of claim 224, wherein said enzymatic nucleic acid molecule comprises any of the NCH ribozyme sequences shown in Table 34.

15 227. The enzymatic nucleic acid molecule of claim 224, wherein said enzymatic nucleic acid molecule is used to treat cancer.

228. The enzymatic nucleic acid molecule of claim 224, wherein said cancer is breast cancer.

20 229. The enzymatic nucleic acid molecule of claim 224, wherein said enzymatic nucleic acid molecule is used to treat conditions associated with the level of HER2 gene.

230. An enzymatic nucleic acid molecule, wherein said enzymatic nucleic acid molecule comprises any of sequence shown as NCH ribozyme sequence in Table 31.

25 231. The enzymatic nucleic acid molecule of claim 224, wherein said enzymatic nucleic acid molecule comprises a substrate binding region which has between 5 and 30 nucleotides complementary to the RNA.

30 232. The enzymatic nucleic acid molecule of claim 224, wherein said enzymatic nucleic acid molecule comprises a substrate binding region which has between 7 and 12 nucleotides complementary to the RNA.

233. A mammalian cell including the enzymatic nucleic acid molecule of claim 224, wherein said mammalian cell is not a living human.
234. The mammalian cell of claim 233, wherein said mammalian cell is a human cell.
- 5 235. A mammalian cell including the enzymatic nucleic acid molecule of claims 186 or 187, wherein said mammalian cell is not a living human.
236. The mammalian cell of claim 235, wherein said mammalian cell is a human cell.
- 10 237. A method for inhibiting expression of HER2 gene in a cell, comprising the step of administering to said cell the enzymatic nucleic acid molecule of claim 224 under conditions suitable for said inhibition.
238. A method of cleaving RNA derived from HER2 gene comprising, contacting the enzymatic nucleic acid molecule of claim 224 with said RNA molecule under conditions suitable for the cleavage of said RNA molecule.
- 15 239. A pharmaceutical composition comprising the enzymatic nucleic acid molecule of any of claims 186 or 187.
240. A pharmaceutical composition comprising the enzymatic nucleic acid molecule of claim 224.
- 20 241. A method of treatment of a patient having a condition associated with the level of HER2, wherein said patient is administered the enzymatic nucleic acid molecule of claim 224 under conditions suitable for said treatment.
242. The method of claim 241, wherein said method is performed in conjunction with one or more other therapies.
- 25 243. The enzymatic nucleic acid molecule of claim 227, wherein said enzymatic nucleic acid molecule is used in conjunction with one or more other therapies.
244. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said nucleic acid molecule comprises at least five ribose residues; a 2'-C-allyl modification at position No. 4 of said enzymatic nucleic acid; at least ten 2'-O-alkyl modifications, and a 3'-cap structure.

245. The enzymatic nucleic acid molecule of claim 244, wherein said 2'-O-alkyl modifications is selected from the group consisting of 2'-O-methyl and 2'-O-allyl.
246. The enzymatic nucleic acid molecule of claim 244, wherein said 3'-cap is 3'-3' inverted abasic moiety.
- 5 247. The enzymatic nucleic acid molecule of claim 244, wherein said 3'-cap is 3'-3' inverted nucleotide.
248. The enzymatic nucleic acid molecule of claim 244, wherein said enzymatic nucleic acid comprises phosphorothioate linkages in at least three of the 5' terminal nucleotides.
- 10 249. The enzymatic nucleic acid molecule of claims 186 or 187, wherein said nucleic acid molecule comprises at least five ribose residues; a 2'-deoxy-2'-amino modification at position Nos. 4 and 7 of said enzymatic nucleic acid; at least ten 2'-O-alkyl modifications, and a 3'- cap structure.
250. The enzymatic nucleic acid molecule of claim 249, wherein said 2'-O-alkyl
15 modifications is selected from the group consisting of 2'-O-methyl and 2'-O-allyl.
251. The enzymatic nucleic acid molecule of claim 249, wherein said 3'-cap is 3'-3' inverted abasic moiety.
252. The enzymatic nucleic acid molecule of claim 249, wherein said 3'-cap is 3'-3' inverted nucleotide.
- 20 253. The enzymatic nucleic acid molecule of claim 249, wherein said enzymatic nucleic acid comprises phosphorothioate linkages in at least three of the 5' terminal nucleotides.
254. The enzymatic nucleic acid molecule of claim 224, wherein said enzymatic nucleic acid molecule comprises at least one sugar modification.
- 25 255. The enzymatic nucleic acid molecule of claim 224, wherein said enzymatic nucleic acid molecule comprises at least one nucleic acid base modification.
256. The enzymatic nucleic acid molecule of claim 224, wherein said enzymatic nucleic acid molecule comprises at least one phosphate backbone modification.

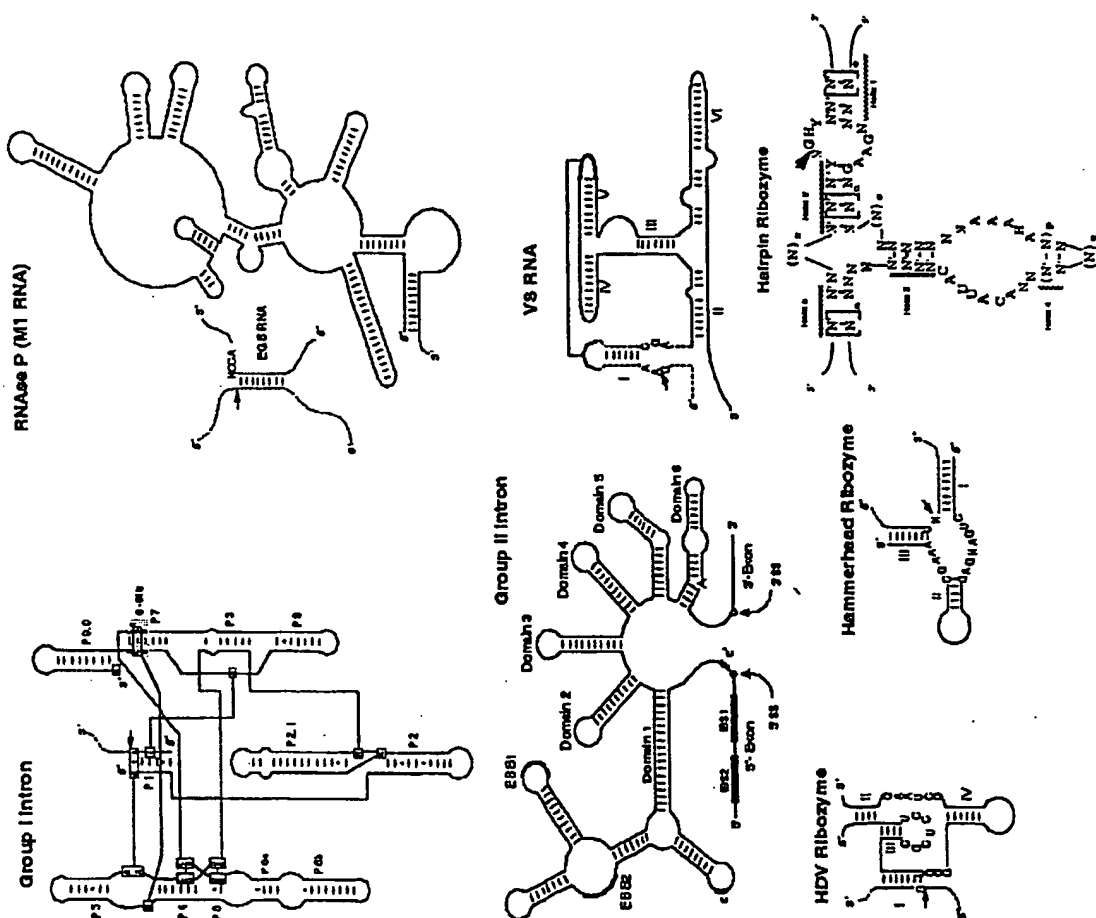
257. The enzymatic nucleic acid molecule of claim 224, wherein said phosphate backbone modification is selected from the group consisting of phosphorothioate, phosphorodithioate and amide.
258. The enzymatic nucleic acid molecule of claim 224, wherein said nucleic acid molecule comprises at least five ribose residues; a 2'-C-allyl modification at position No. 4 of said enzymatic nucleic acid; at least ten 2'-O-alkyl modifications, and a 3'-cap structure.
259. The enzymatic nucleic acid molecule of claim 258, wherein said 2'-O-alkyl modifications is selected from the group consisting of 2'-O-methyl and 2'-O-allyl.
260. The enzymatic nucleic acid molecule of claim 258, wherein said 3'-cap is 3'-3' inverted abasic moiety.
261. The enzymatic nucleic acid molecule of claim 258, wherein said 3'-cap is 3'-3' inverted nucleotide.
262. The enzymatic nucleic acid molecule of claim 258, wherein said enzymatic nucleic acid comprises phosphorothioate linkages in at least three of the 5' terminal nucleotides.
263. The enzymatic nucleic acid molecule of claim 224, wherein said nucleic acid molecule comprises at least five ribose residues; a 2'-deoxy-2'-amino modification at position Nos. 4 and 7 of said enzymatic nucleic acid; at least ten 2'-O-alkyl modifications, and a 3'-cap structure.
264. The enzymatic nucleic acid molecule of claim 263, wherein said 2'-O-alkyl modifications is selected from the group consisting of 2'-O-methyl and 2'-O-allyl.
265. The enzymatic nucleic acid molecule of claim 263, wherein said 3'-cap is 3'-3' inverted abasic moiety.
266. The enzymatic nucleic acid molecule of claim 263, wherein said 3'-cap is 3'-3' inverted nucleotide.
267. The enzymatic nucleic acid molecule of claim 263, wherein said enzymatic nucleic acid comprises phosphorothioate linkages in at least three of the 5' terminal nucleotides.

268. The enzymatic nucleic acid molecule of claim 186, wherein said enzymatic nucleic acid molecule is capable of down-regulating the expression of protein kinase C alpha (PKC alpha) gene.
269. A method for inhibiting expression of a PKC alpha gene in a cell, comprising
5 the step of administering to said cell the enzymatic nucleic acid molecule of claim 268 under conditions suitable for said inhibition.
270. A method of cleaving a PKC alpha RNA molecule comprising, contacting the enzymatic nucleic acid molecule of claim 268 with said separate PKC alpha RNA molecule under conditions suitable for the cleavage of said PKC alpha RNA
10 molecule.
271. The method of claim 270, wherein said cleavage is carried out in the presence of a divalent cation.
272. The method of claim 271, wherein said divalent cation is Mg^{2+} .
273. The enzymatic nucleic acid molecule of claim 268, wherein said enzymatic
15 nucleic acid molecule has an endonuclease activity to cleave RNA derived from PKC alpha gene.
274. The enzymatic nucleic acid molecule of claim 273, wherein said enzymatic nucleic acid molecule comprises sequences complementary to any of NCH substrate sequence of Table 63.
- 20 275. The enzymatic nucleic acid molecule of claim 273 wherein said enzymatic nucleic acid molecule comprises any of the NCH ribozyme sequences shown in Table 63.
276. The enzymatic nucleic acid molecule of claim 268, wherein said enzymatic nucleic acid molecule is used to treat cancer.
- 25 277. The enzymatic nucleic acid molecule of claim 276, wherein said cancer is selected from the group consisting of lung, breast, colon, prostate, bladder, ovary, melanoma, and glioblastoma cancer.
278. The enzymatic nucleic acid molecule of claim 268, wherein said enzymatic
30 nucleic acid molecule is used to treat conditions associated with the level of PKC alpha gene.

279. The enzymatic nucleic acid molecule of claim 268, wherein said D and E independently has between 5 and 30 nucleotides complementary to the RNA.
280. The enzymatic nucleic acid molecule of claim 268, wherein said D and E independently has between 7 and 12 nucleotides complementary to the RNA.
- 5 281. A mammalian cell including the enzymatic nucleic acid molecule of claim 268, wherein said mammalian cell is not a living human.
282. The mammalian cell of claim 281, wherein said mammalian cell is a human cell.
- 10 283. A pharmaceutical composition comprising the enzymatic nucleic acid molecule of claim 238.
284. A pharmaceutical composition comprising the enzymatic nucleic acid molecule of claim 273.
- 15 285. A method of treatment of a patient having a condition associated with the level of PKC alpha, wherein said patient is administered the enzymatic nucleic acid molecule of claim 268 under conditions suitable for said treatment.
286. The method of claim 285, wherein said method is performed in conjunction with one or more other therapies.
287. The enzymatic nucleic acid molecule of claim 286, wherein said enzymatic nucleic acid molecule is used in conjunction with one or more other therapies.
- 20 288. An antisense nucleic acid molecule comprising sequence complementary to any of substrate sequence in Tables 13-23.
289. The antisense nucleic acid molecule of claim 288, wherein said enzymatic nucleic acid is chemically synthesized.
- 25 290. The antisense nucleic acid molecule of claim 288, wherein said antisense nucleic acid comprises at least one 2'-sugar modification.
291. The antisense nucleic acid molecule of claim 288, wherein said antisense nucleic acid comprises at least one nucleic acid base modification.
292. The antisense nucleic acid molecule of claim 288, wherein said antisense nucleic acid comprises at least one phosphate backbone modification.

293. A mammalian cell including the antisense nucleic acid molecule of claim 288, wherein said mammalian cell is not a living human.
294. The mammalian cell of claim 293, wherein said mammalian cell is a human cell.

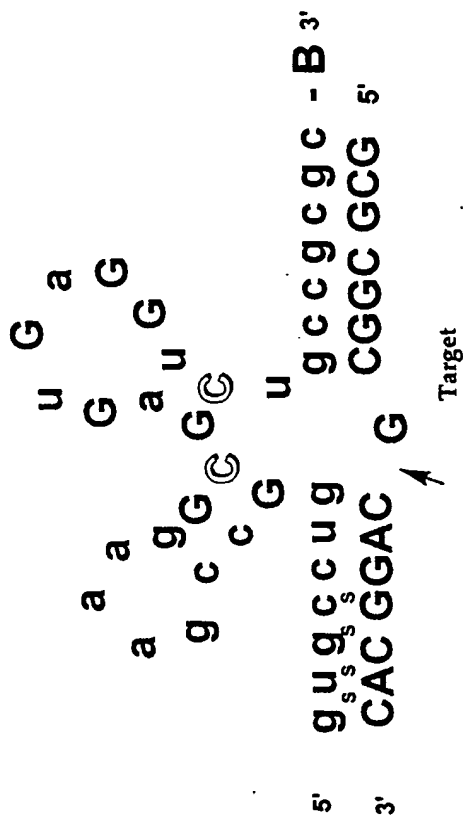
Figure 1: Ribozyme Motifs



[illegible]

Figure 4: Stabilized Zinzyme Ribozyme Motif

Zinzyme A-motif RZ



Legend

Uppercase Indicates natural ribo residues

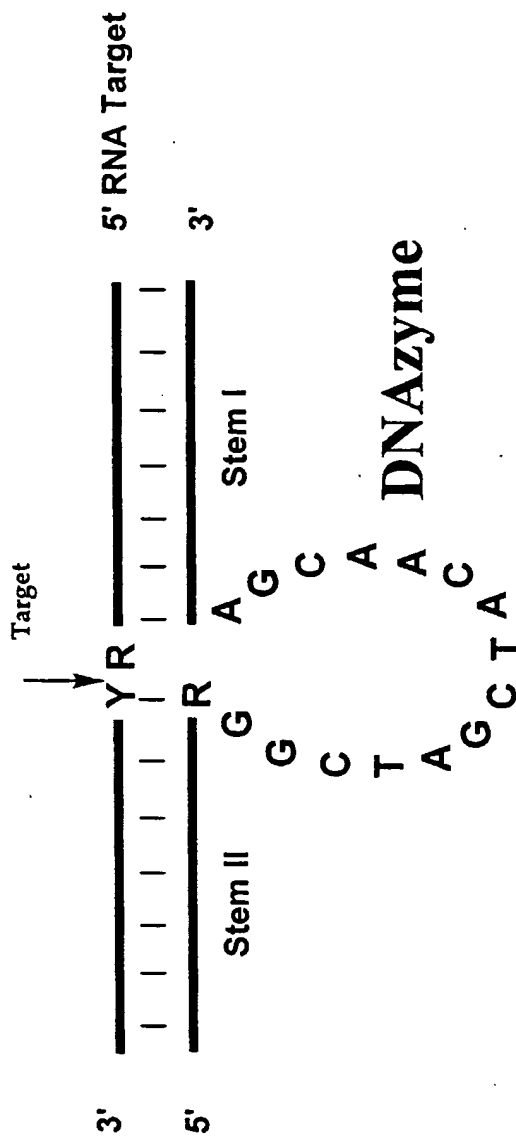
Ⓒ indicates 2' - d-NH₂-C

Lowercase: 2'-O- Me

Subscript _s Indicates phosphothioate linkage

B: 3'-3' abasic moiety

Figure 5: DNAzyme Motif

Legend

Y = U or C

R = A or G

Figure 7: Examples of Nuclease Stable Ribozyme Motifs

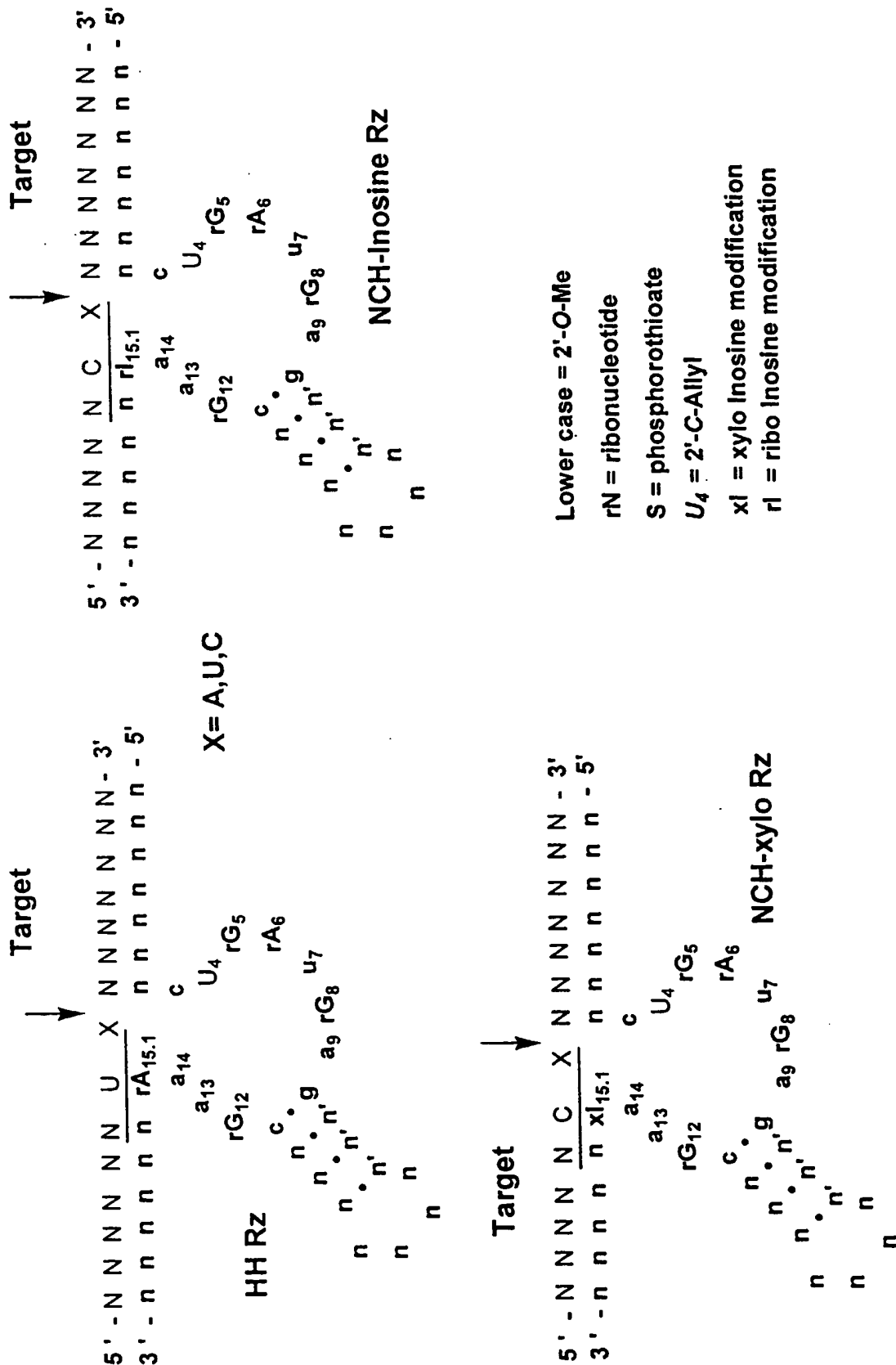


Figure 8: Inhibition of Cell Proliferation by Anti-Her2 Ribozymes

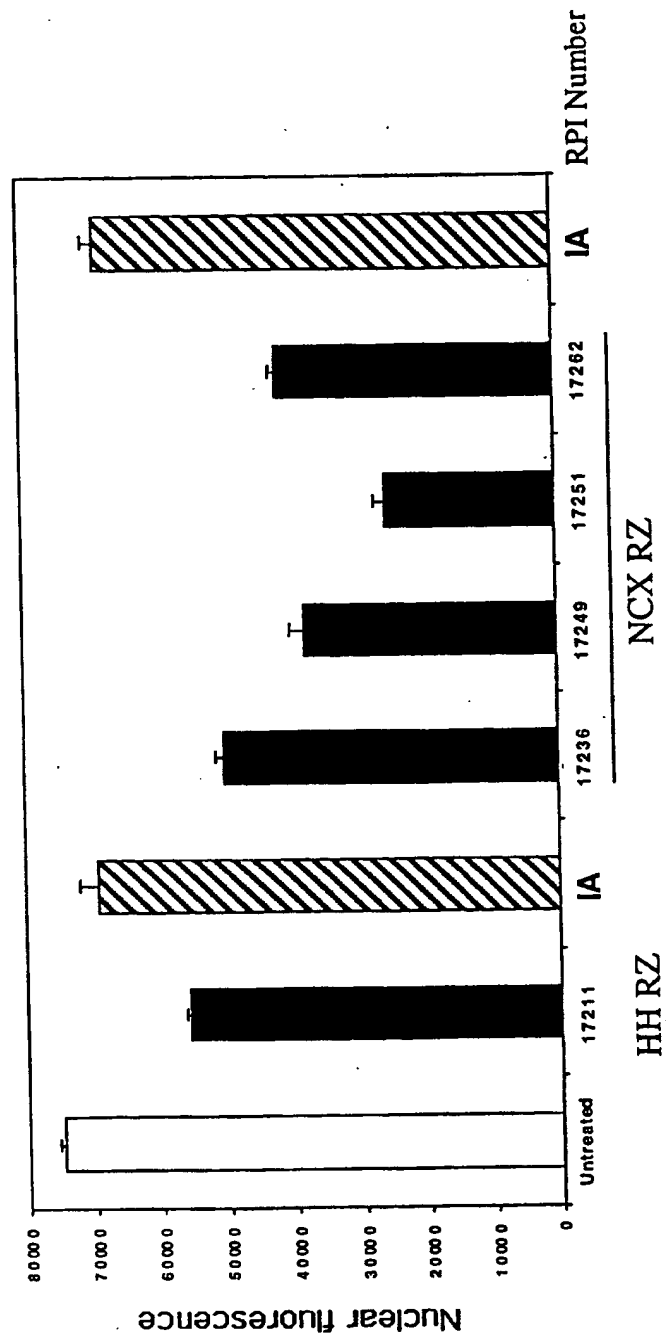
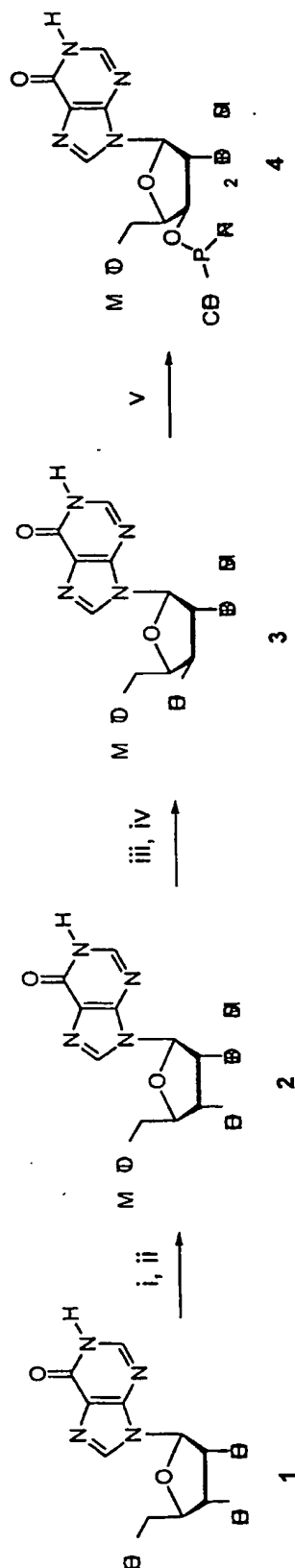


Figure 9: Synthesis of b-D-xylofuranosyl hypoxanthine 3'-phosphoramidite

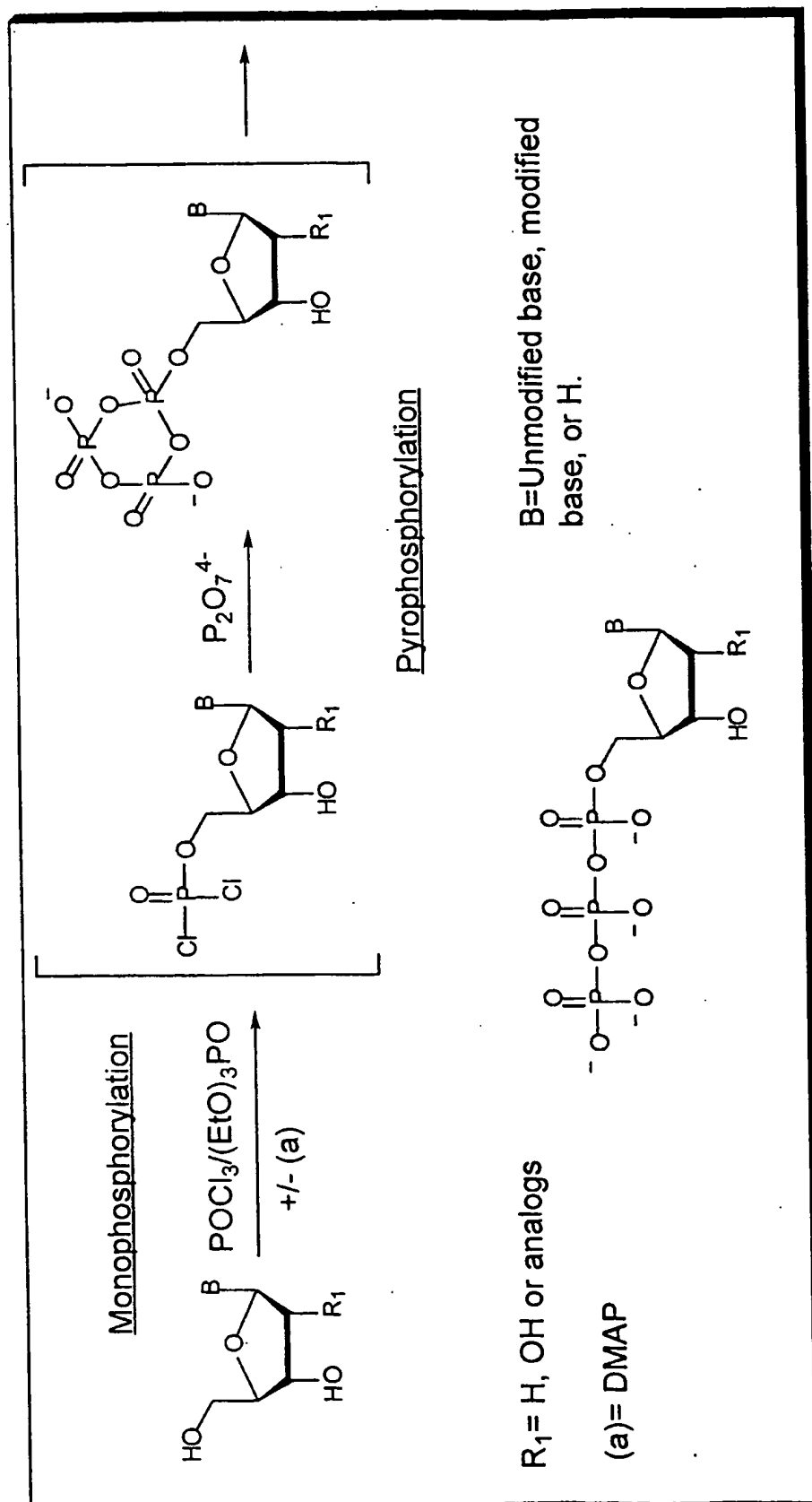


MMT = 4-methoxytriphenylmethyl

TBDMS = *t*-butyldimethylsilyl

Reagents and Conditions: (i) MMT-Cl/Pyr-DMSO, rt, 48 h; (ii) TBDMS-Cl/AgNO₃/Pyr/THF; (iii) CrO₃/Pyr/Ac₂O/DCM, rt, 1 h; (iv) NaB(OAc)₃/EtOH, rt, overnight; (v) 2-Cyanoethyl-*N,N*-diisopropylchlorophosphoramidite/1-Melm/DIPEA/DCM, rt, 2 h.

Figure 10: One-Pot Formation of Nucleoside-5'-triphosphates



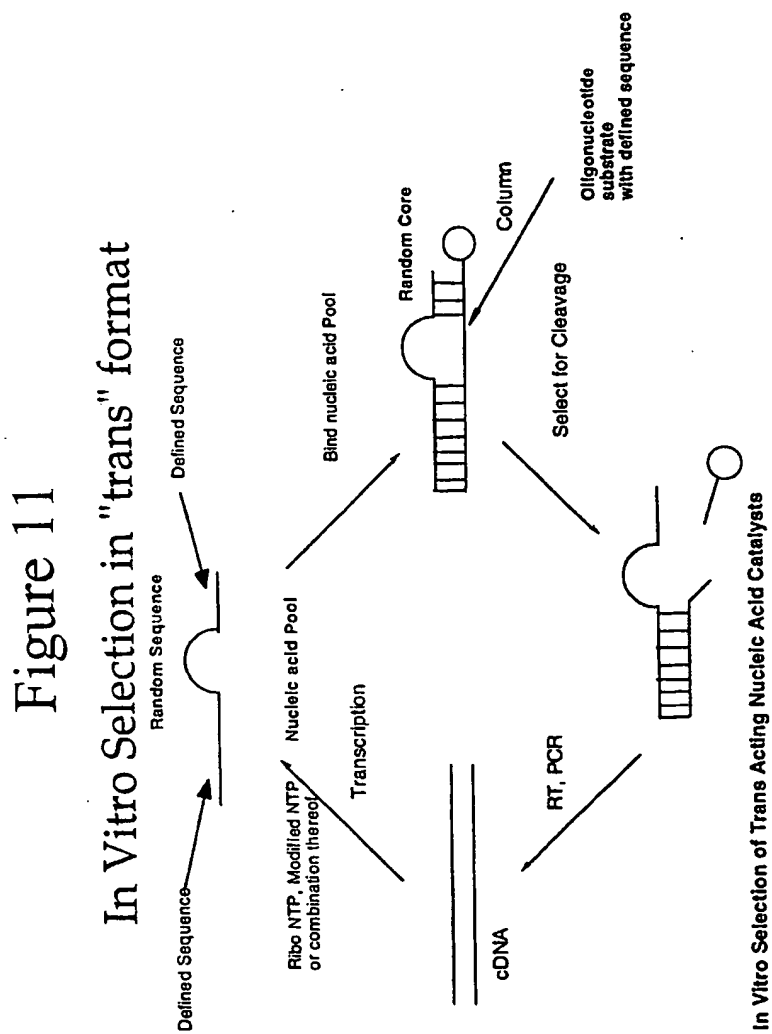


Figure 12. Removal of "parasitic RNA" using a Second Selection column

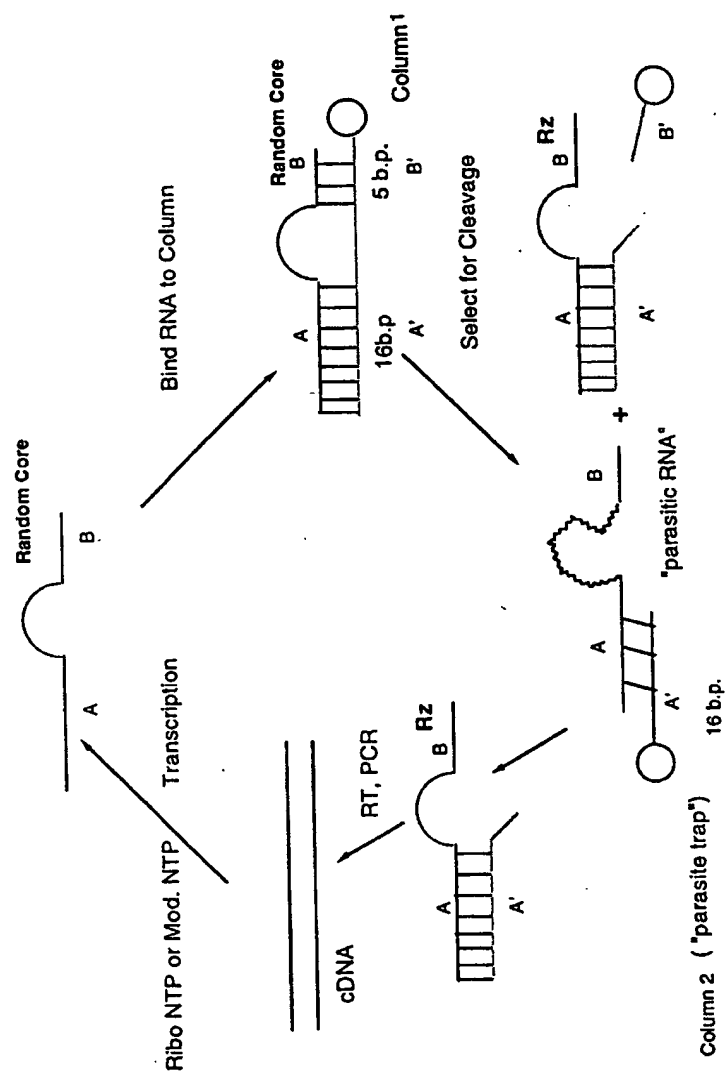


FIGURE 14. Dual Reporter System for Cytoplasmic HCV Target

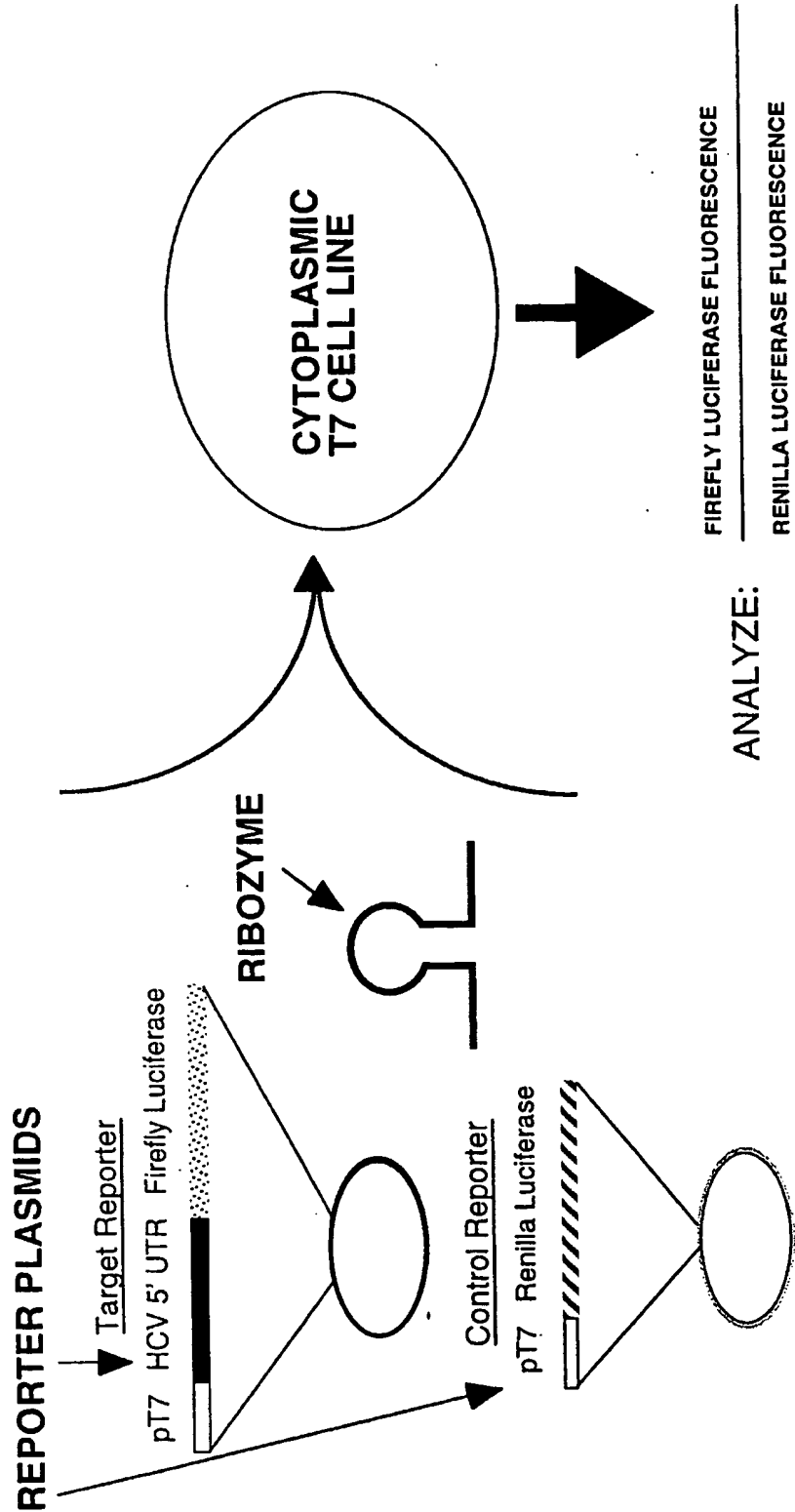


Figure 15. Dose-dependent inhibition of HCV-IRES mediated luciferase activity

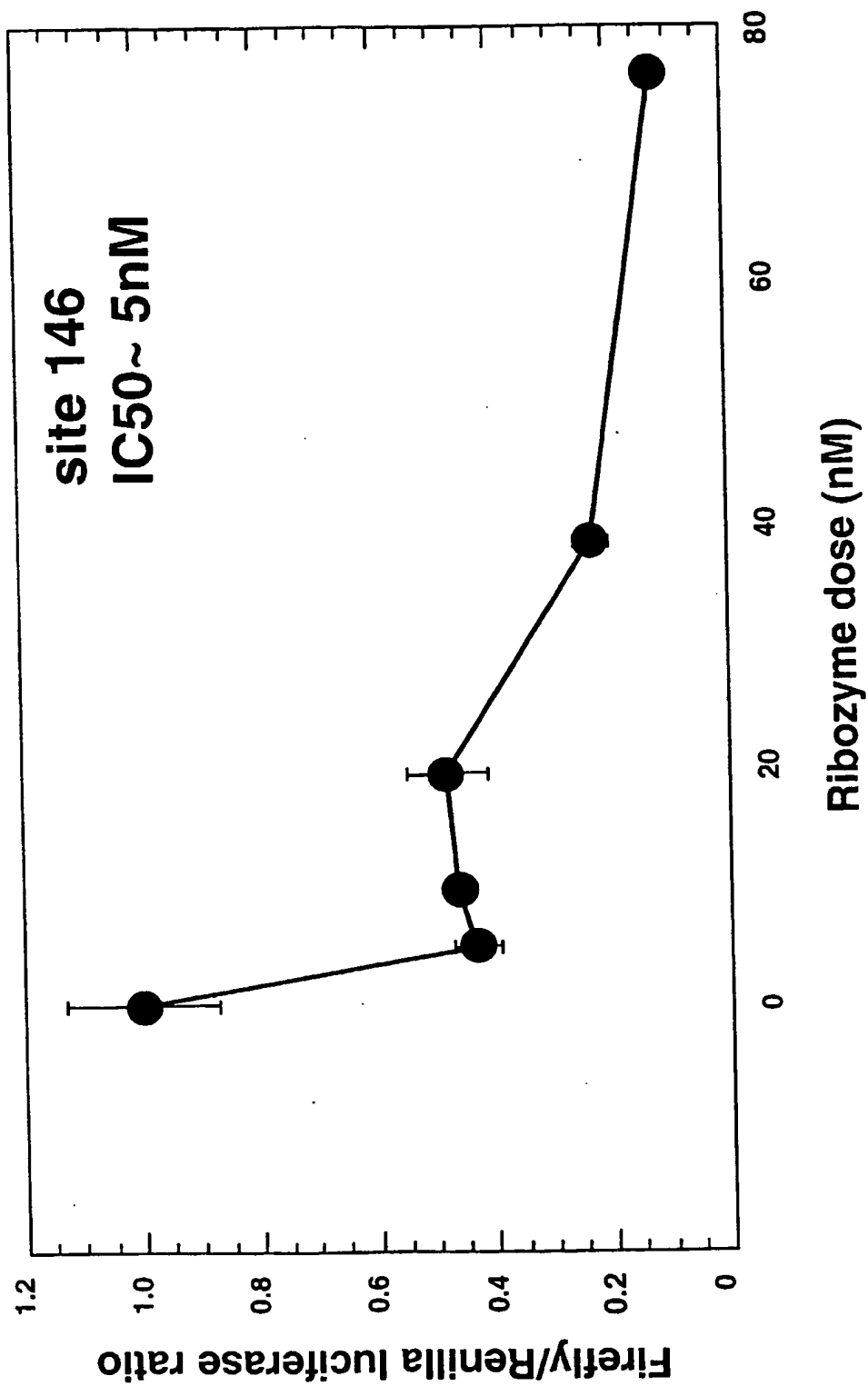
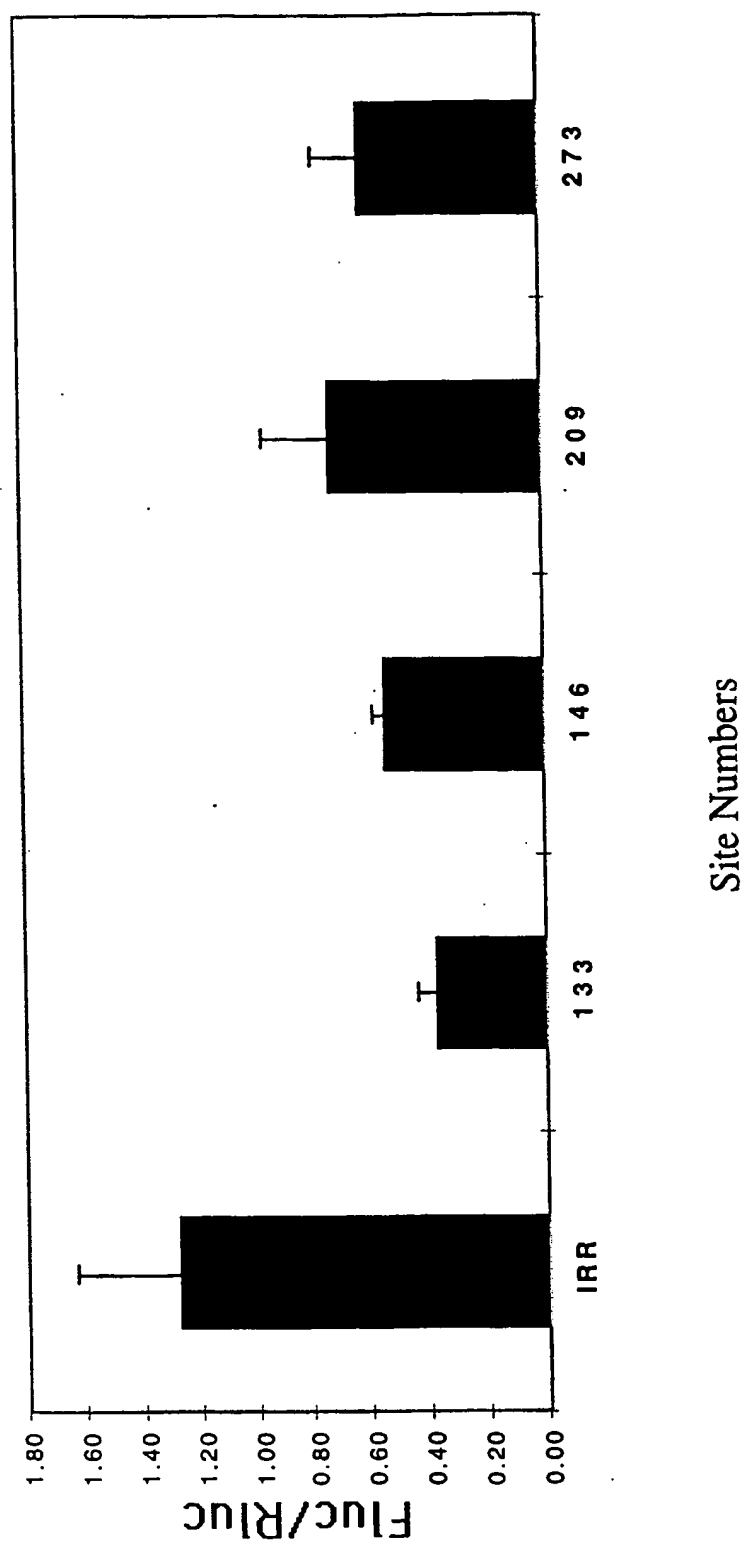


Figure 16. Efficacious Ribozymes Targeting 5'UTR HCV RNA



Sequence and chemical compositions for site numbers are given in table XII

Figure 17. Characterized Class II Enzymatic Nucleic Acid Motifs

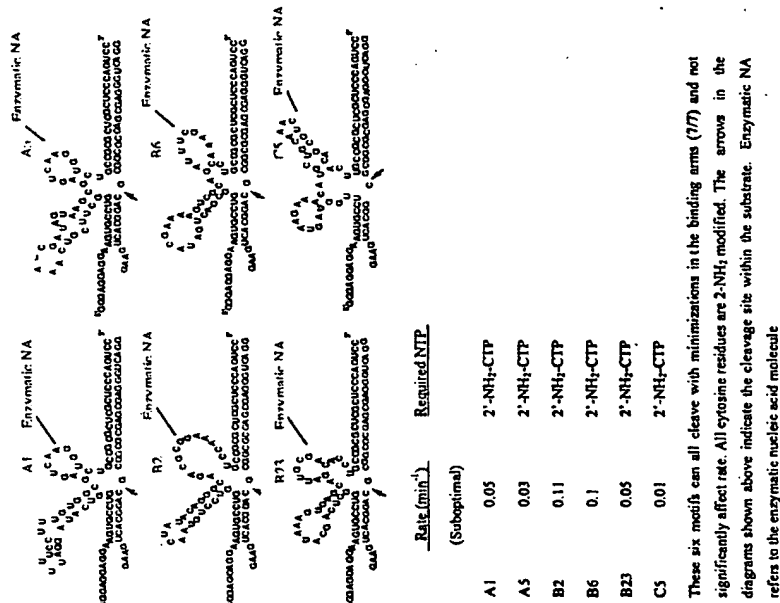
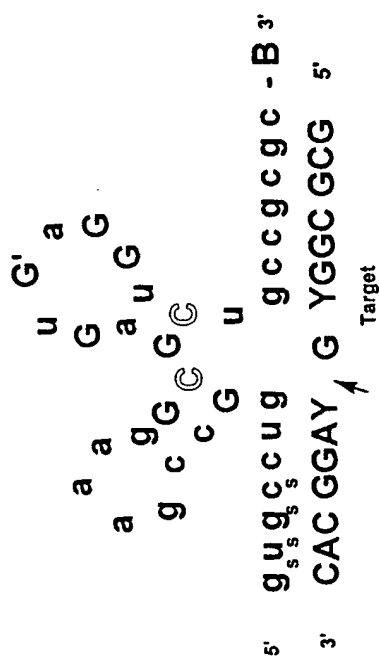


Figure 18: Chemically Stabilized Class II Motif

**Legend**

Uppercase indicates natural ribo residues

C indicates 2' - d-NH₂-C

Lowercase: 2'-O- Me

Subscript s indicates phosphothioate linkage

B: 3'-3' abasic moiety

Y = U or C

G' can be G, ca, or caa

The gaaa tetraloop can be replaced by 18 atom polyethylene glycol (Spacer)

All ribo G's can be replaced with 2'-O-methyl/ G

Figure 19: Substrate specificities of Class II (zinzyme) ribozymes

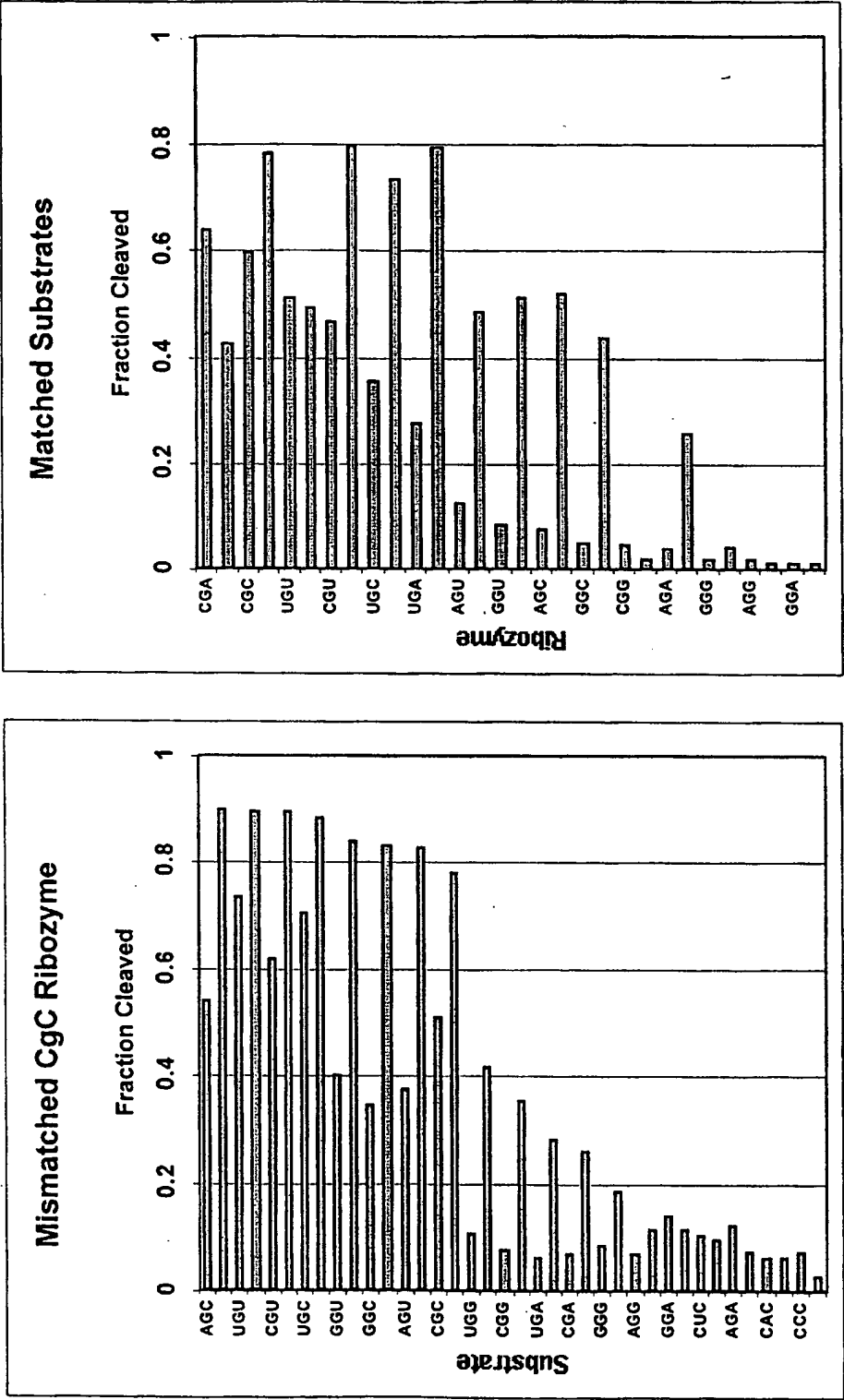
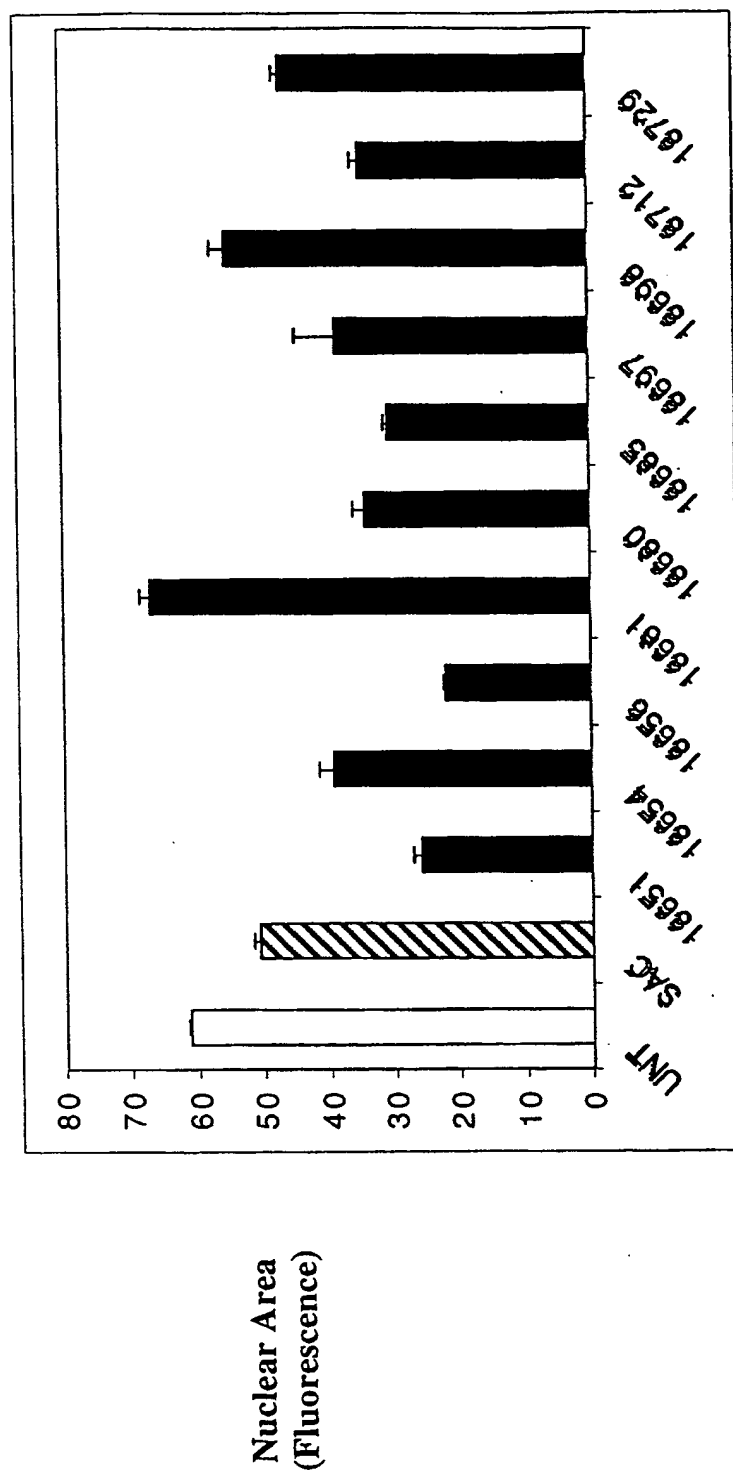
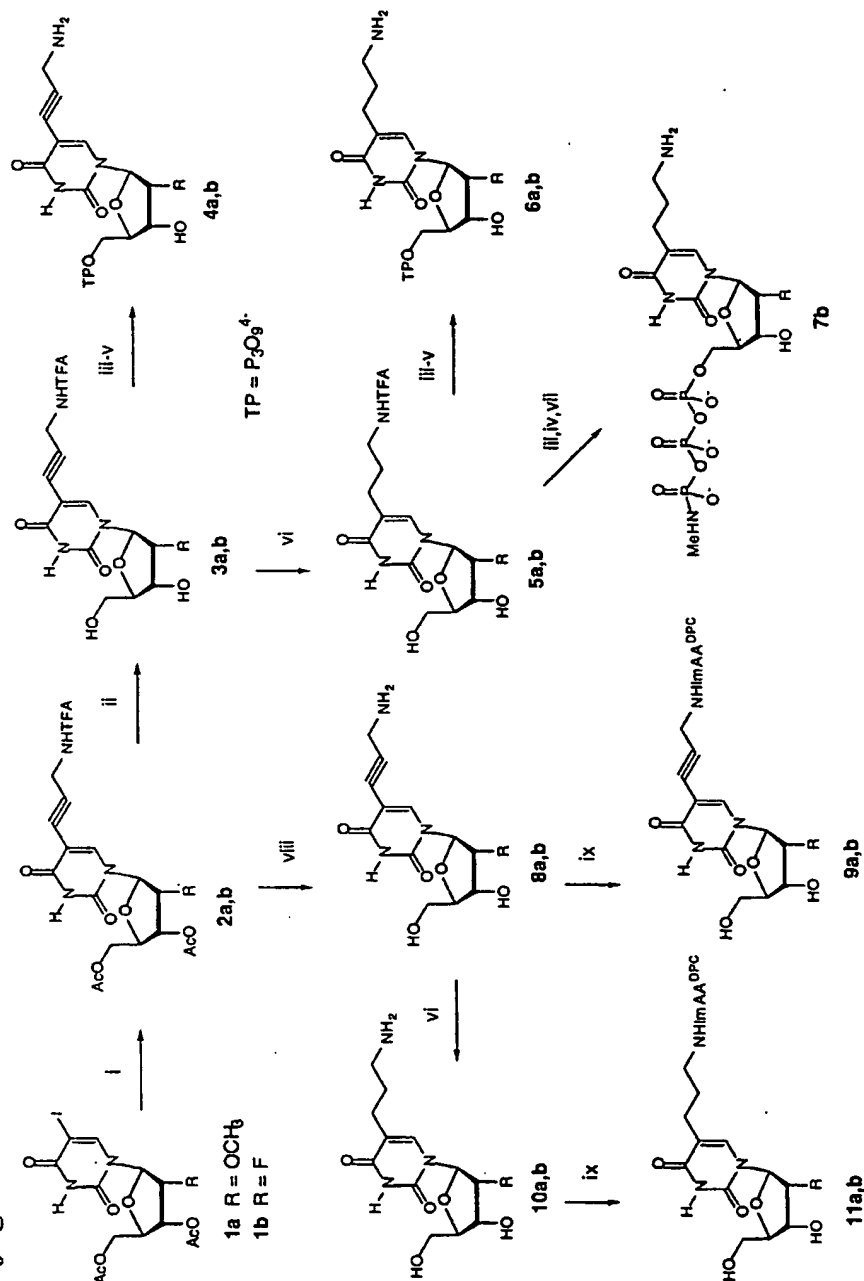


Figure 20: Representative data of HER2 cell proliferation primary screen of Class II (zincyme) Ribozymes



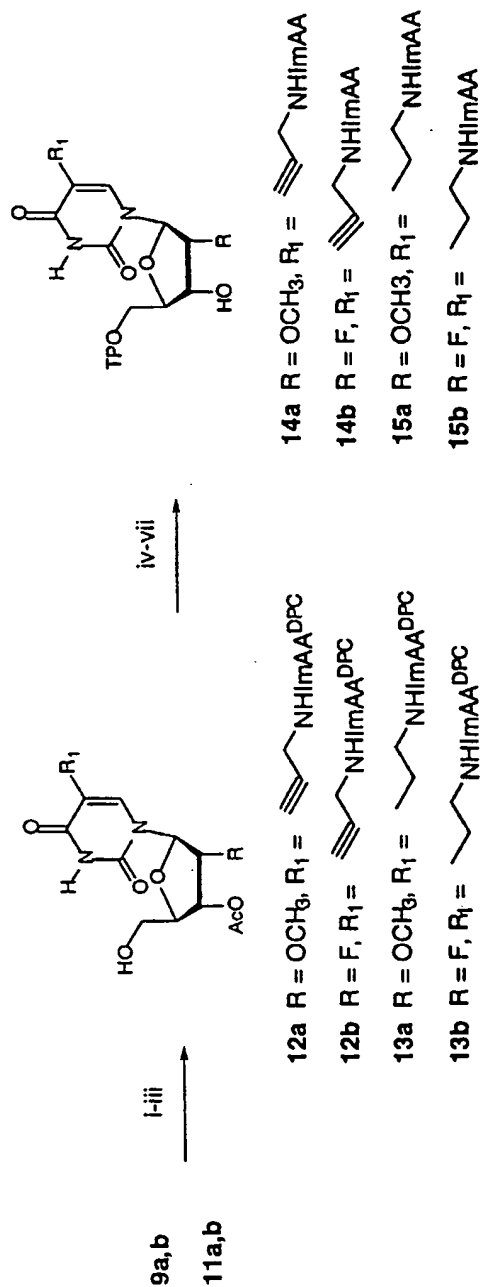
Treatment (RPI number)

SKBR3 breast carcinoma cells
 2 µg/mL RPI9649 (lipid)
 200 nM ribozymes
 120 hour timepoint
 UNT = untreated
 SAC = scrambled attenuated control



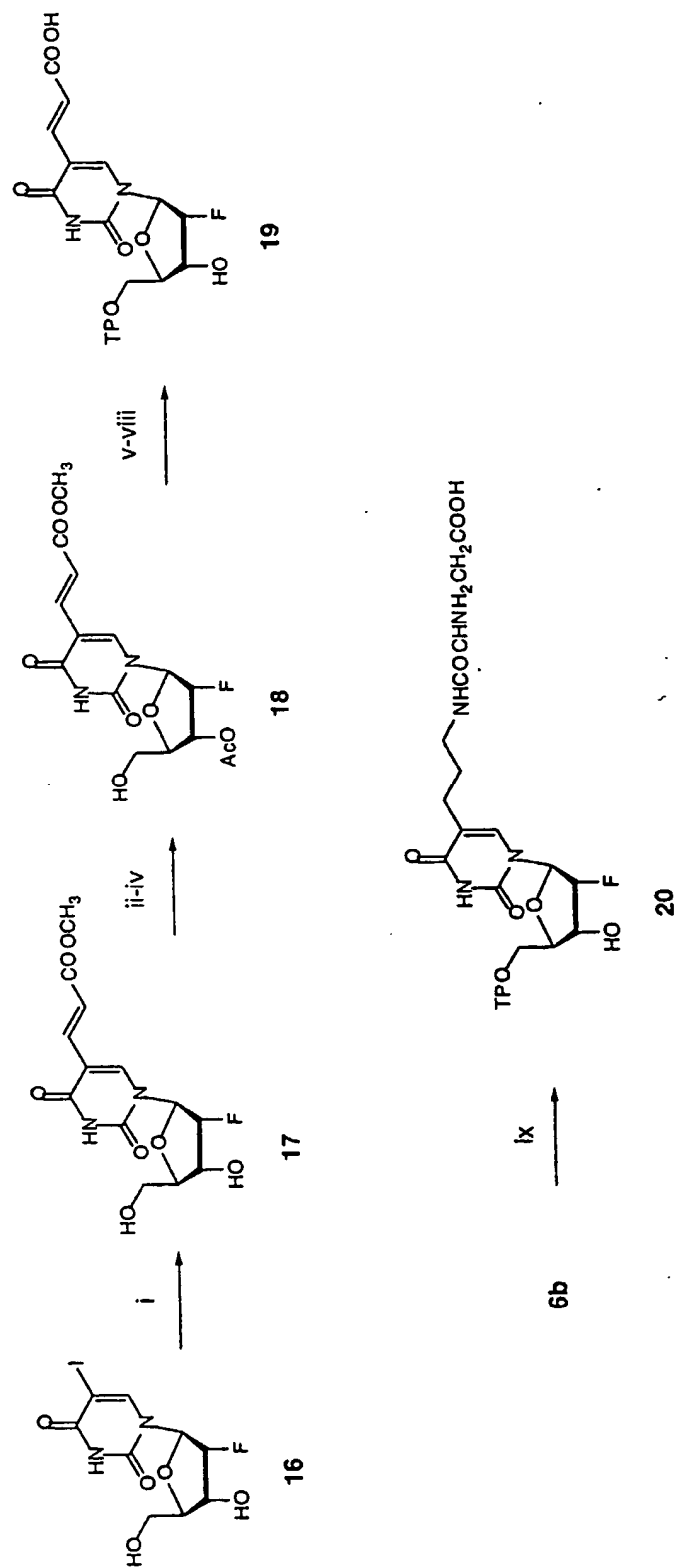
Reagents and Conditions: (I) *N*-TFA propargylamine, CuI, tetrakis(Ph₃P)Pd(0), Et₃N, DMF, 16 h, (ii) aq NaOH, pyr, MeOH, 0 °C, 1 h, (iii) POCl₃, Proton-Sponge, (EtO)₃PO, 2 h, (iv) *n*-Bu₃N PPI, MeCN, 15 min., (v) 1M Et₃NH⁺HCO₃⁻, then NH₄OH, 16 h, (vi) H₂, 5% Pd-C, 24 h, 40 psi, (vii) 40% MeNH₂, 3 h, (viii) NH₄OH, 4 °C, 16 h, (ix) ImAA^{DPC}, EDCHCl, DMF, 16 h.

Figure 22: Synthesis of 5-[3-(N-4-imidazoleacetylaminopropynyl)uridine 5'-triphosphates



Reagents and Conditions: (i) DMT-Cl, pyr, 16 h, (ii) Ac₂O, pyr, 2 h, (iii) 3%TCA, CH₂Cl₂, 2 h, (iv) 2-Cl-4H-1,3,2-benzodioxaphosphorin-4-one, pyr, dioxane, 30 min., (v) *n*-Bu₃N PPI, DMF, 30 min., (vi) I₂, pyr-H₂O, 20 min., (vii) NH₄OH, 2 h.

Figure 23: Synthesis of Carboxylate tethered uridine 5'-triphosphates



Reagents and Conditions: (1) methyl acrylate, Ph_3P , $\text{Pd}(\text{II})$ acetate, Et_3N , dioxane, 30 min., reflux, (ii) DMT-Cl , pyr, 16 h. (iii) Ac_2O , pyr, 3 h. (iv) 3% TCA , CH_2Cl_2 , 1 h, (v) 2-Cl-4H-1,3,2-benzodioxaphosphorin-4-one, pyr, dioxane, 30 min., (vi) $n\text{-Bu}_3\text{N}$ PPI, DMF, 30 min., (vii) I_2 , pyr- H_2O , 20 min., (viii) 1N NaOH , 5 h, (ix) Fmoc-Asp-OFm NHS-ester, DMF-0.1M $\text{Na}_2\text{B}_4\text{O}_7$, 16 h, then Et_3NH , 3 h.

Stem-Loop 3

Deleted
Krel = 0.08

Stem-Loop 2

K rel = 0.18

Stem-Loop 4

K rel = 0.5

Stem-Loop 5

K rel = 0.23

Stem-Loop 6

0.78

Cleavage site

U,C = 2'-NH₂-U,C
Bold = ribo A,G
lower case = Substrate

Figure 26: Class I ribozyme Stem truncation and Loop replacement

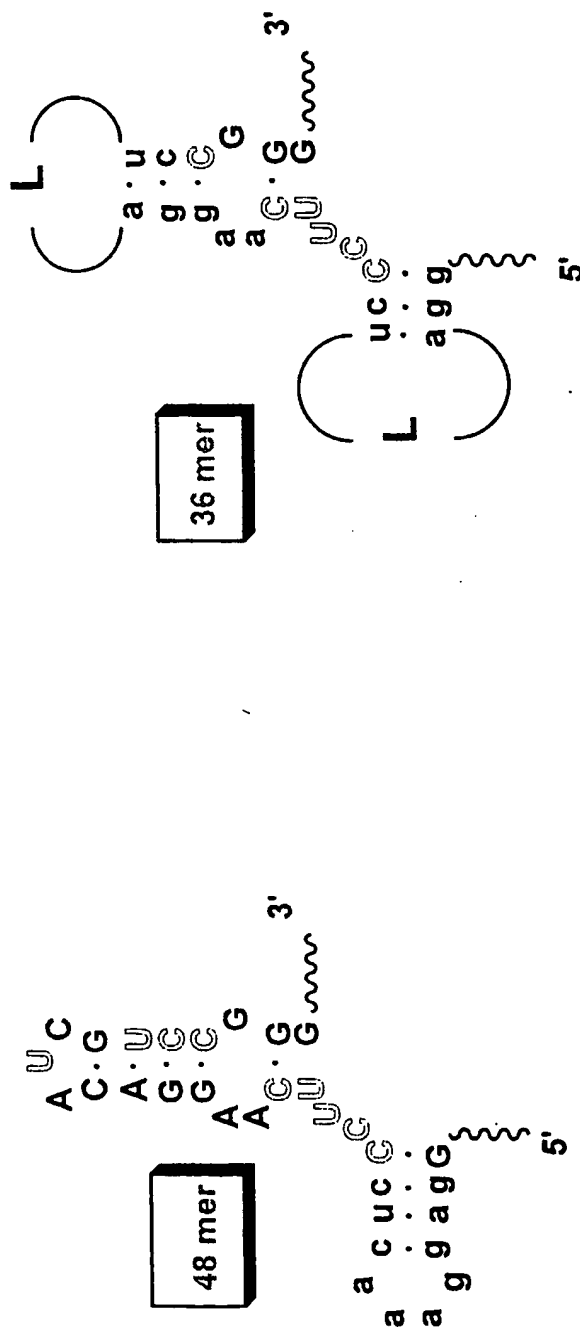
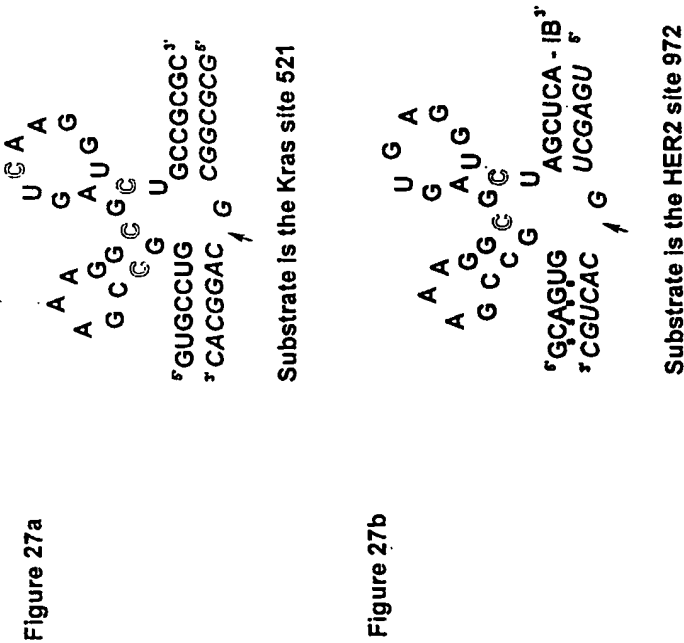


Figure 27: Non ribo Class II (zinzyme) motifs



Legend
Italic indicates natural ribo residues
G indicates 2'-NH₂-C
A G C U indicates 2'-OH residues
Subscript 'a' indicates phosphothioate linkage
IB indicates inverted deoxy abasic residue

Figure 28: Non ribo Class II (zinzyme) cleavage reactions

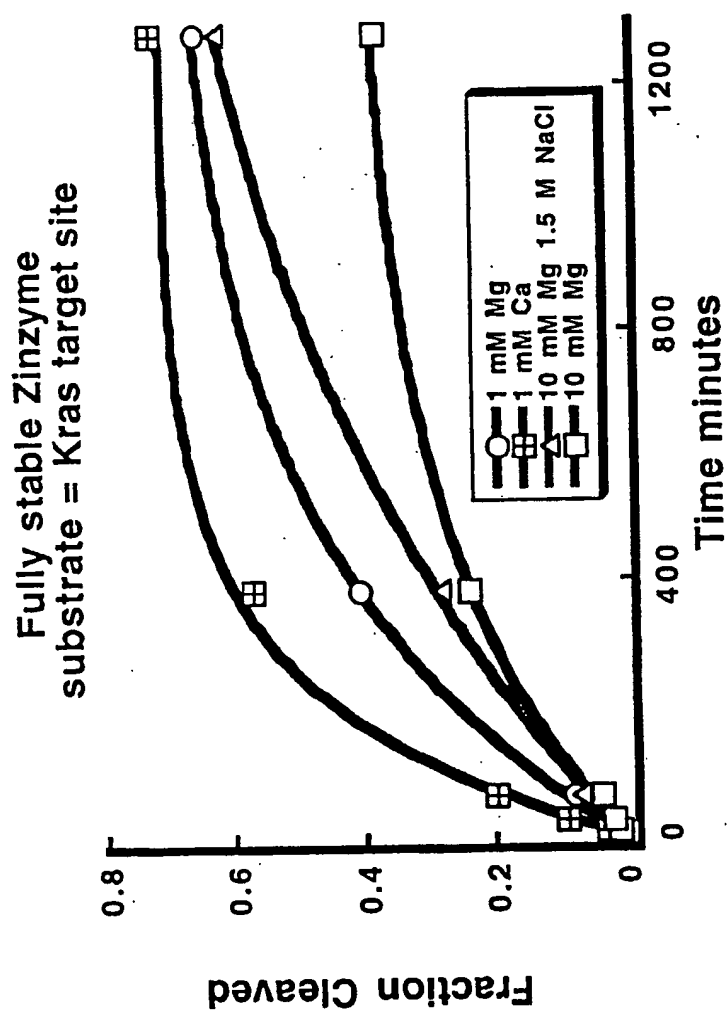
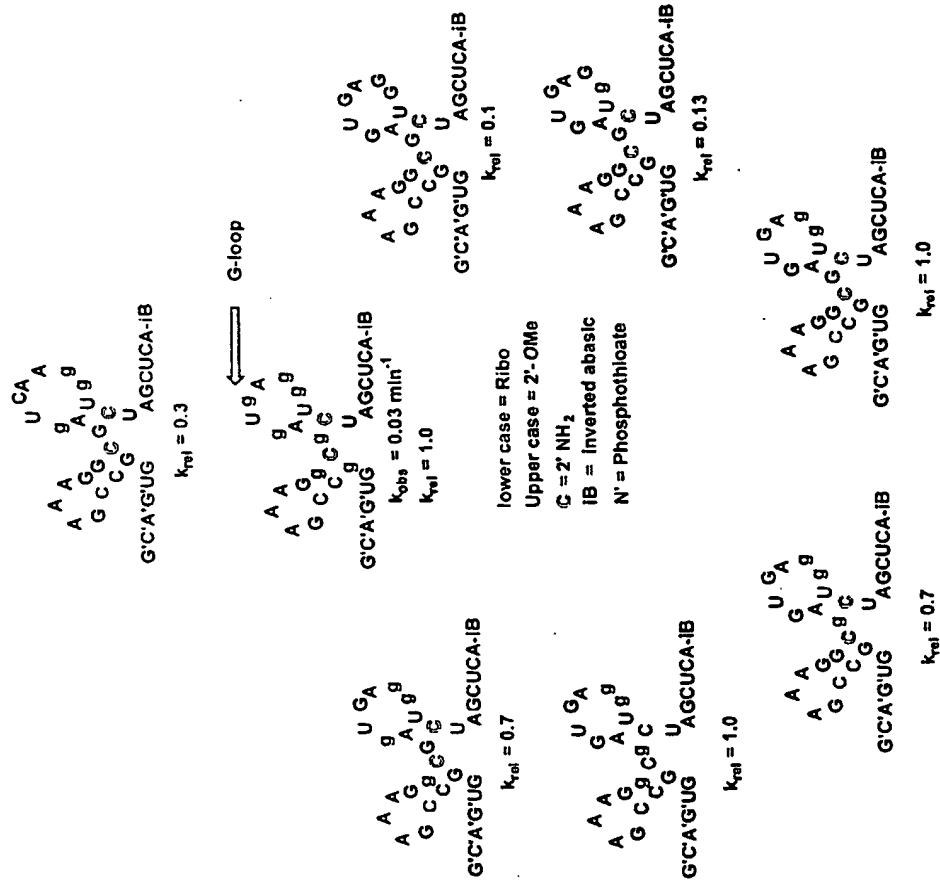


Figure 29: Positional testing of ribose positions in Class II (zinczyme) nucleic acid catalysts targeting HER2 site 972



**All assays done under selection
conditions - physiological buffer
1 mM Mg, 1 mM Ca 37° C
Substrate 15-mer HER2 site 972**

Figure 30: RPI 18656 Mediated Decrease in HER2 RNA
site 972 vs SAC

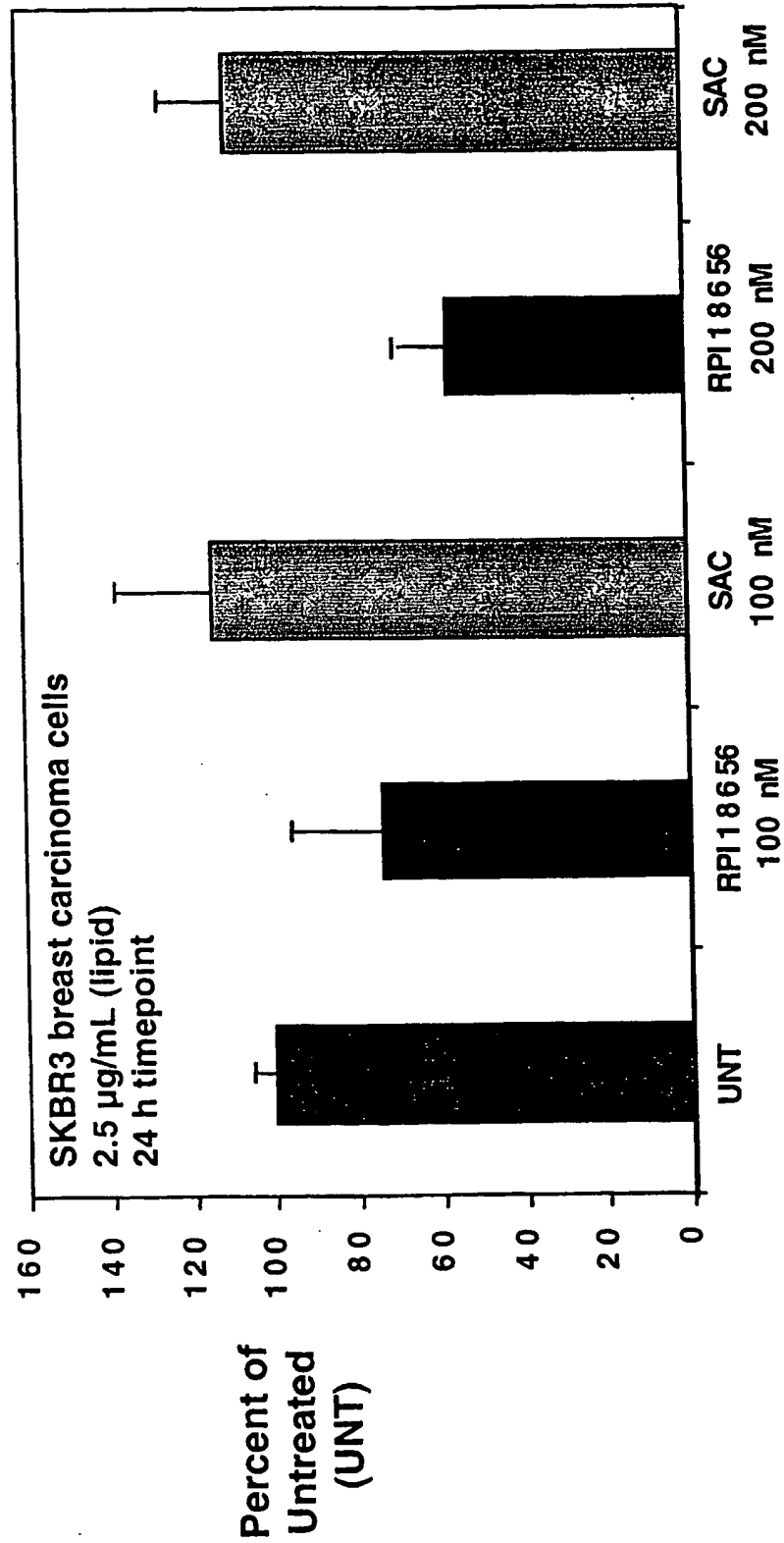


Figure 31: Dose Response of RPI 18656 Against Site 972 in Antiproliferation Assay

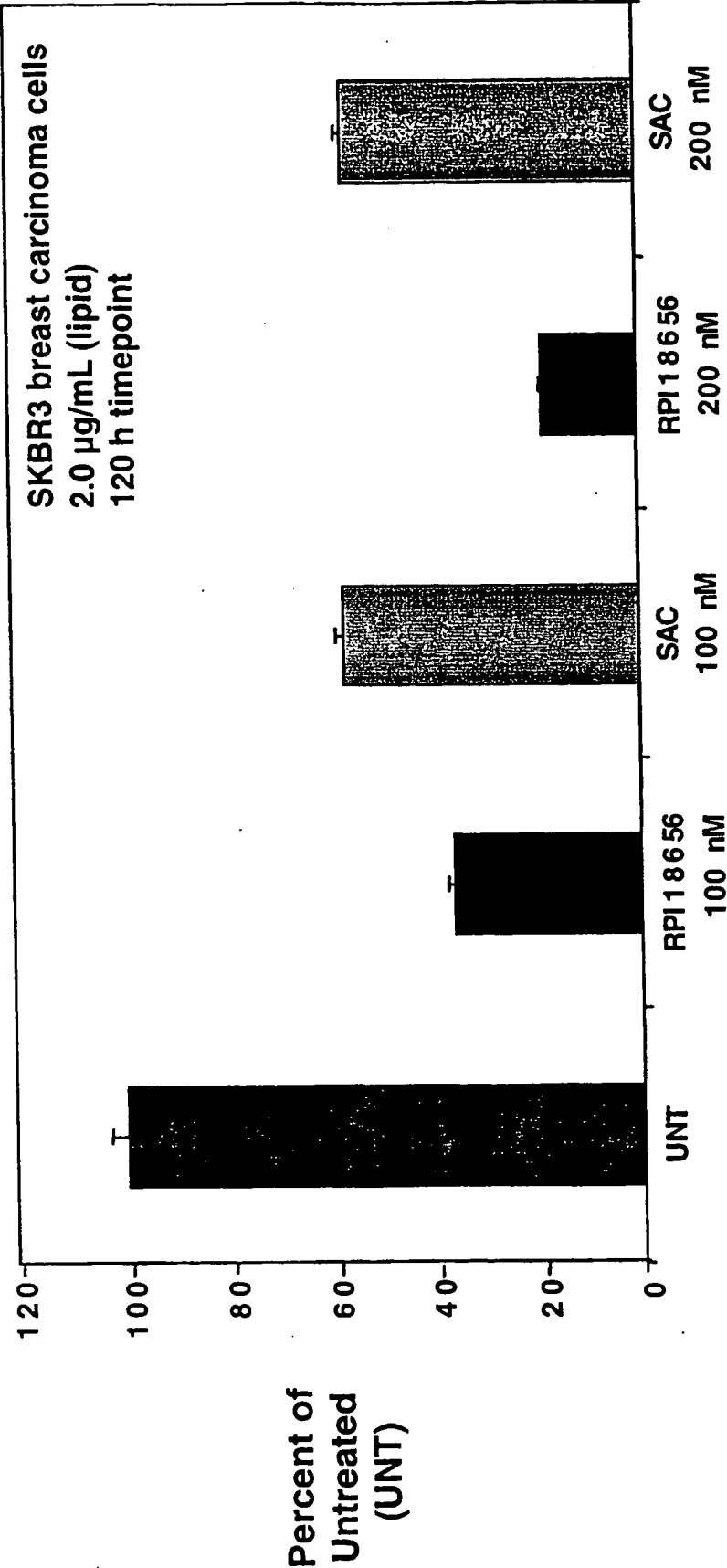


Figure 32: Dose-Dependent HER2 RNA Reduction after
Treatment with RPI 19293

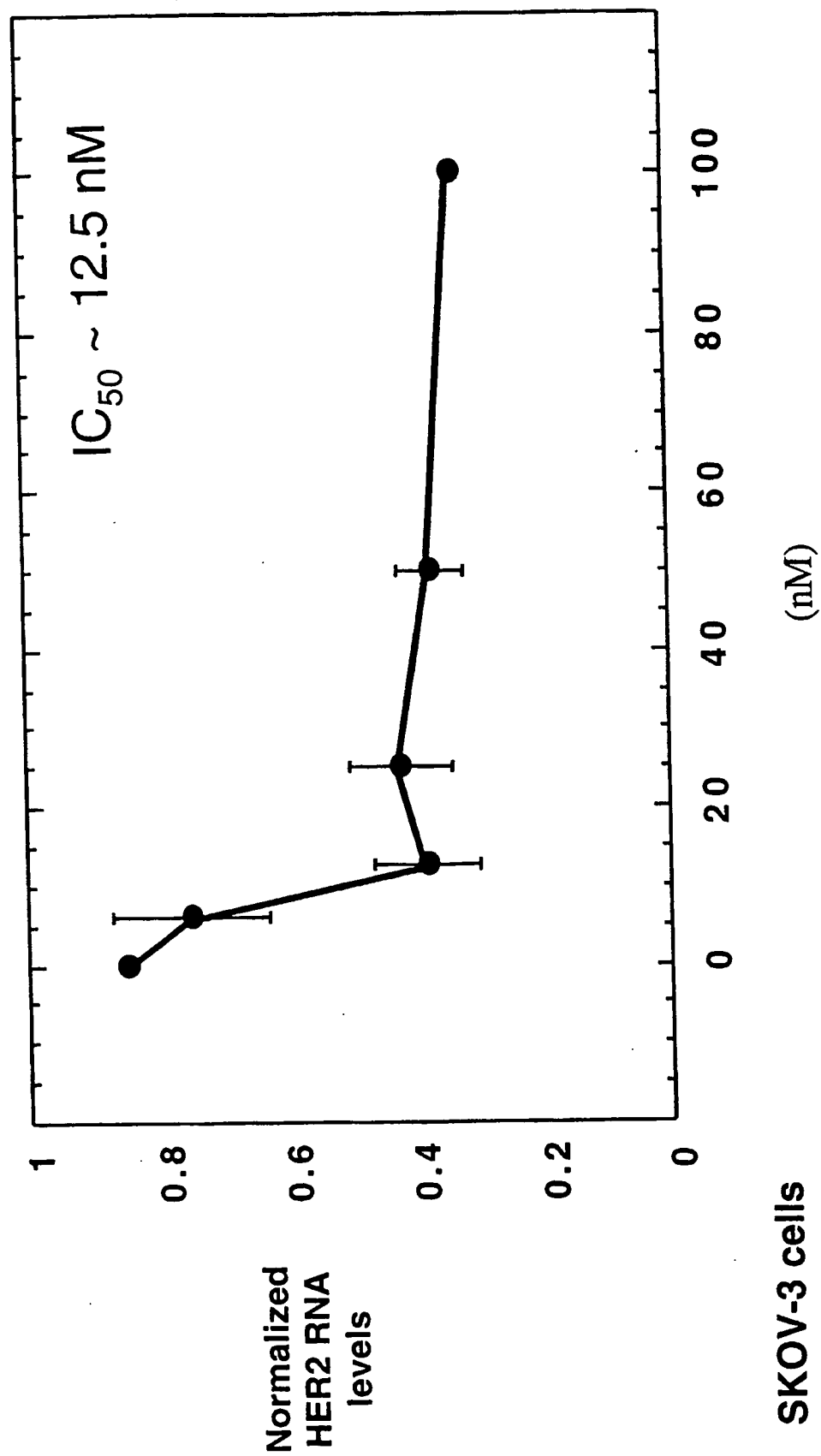


Figure 33: Dose-Dependent HER2 RNA Reduction & Inhibition of Cell Proliferation
(RPI.19293)

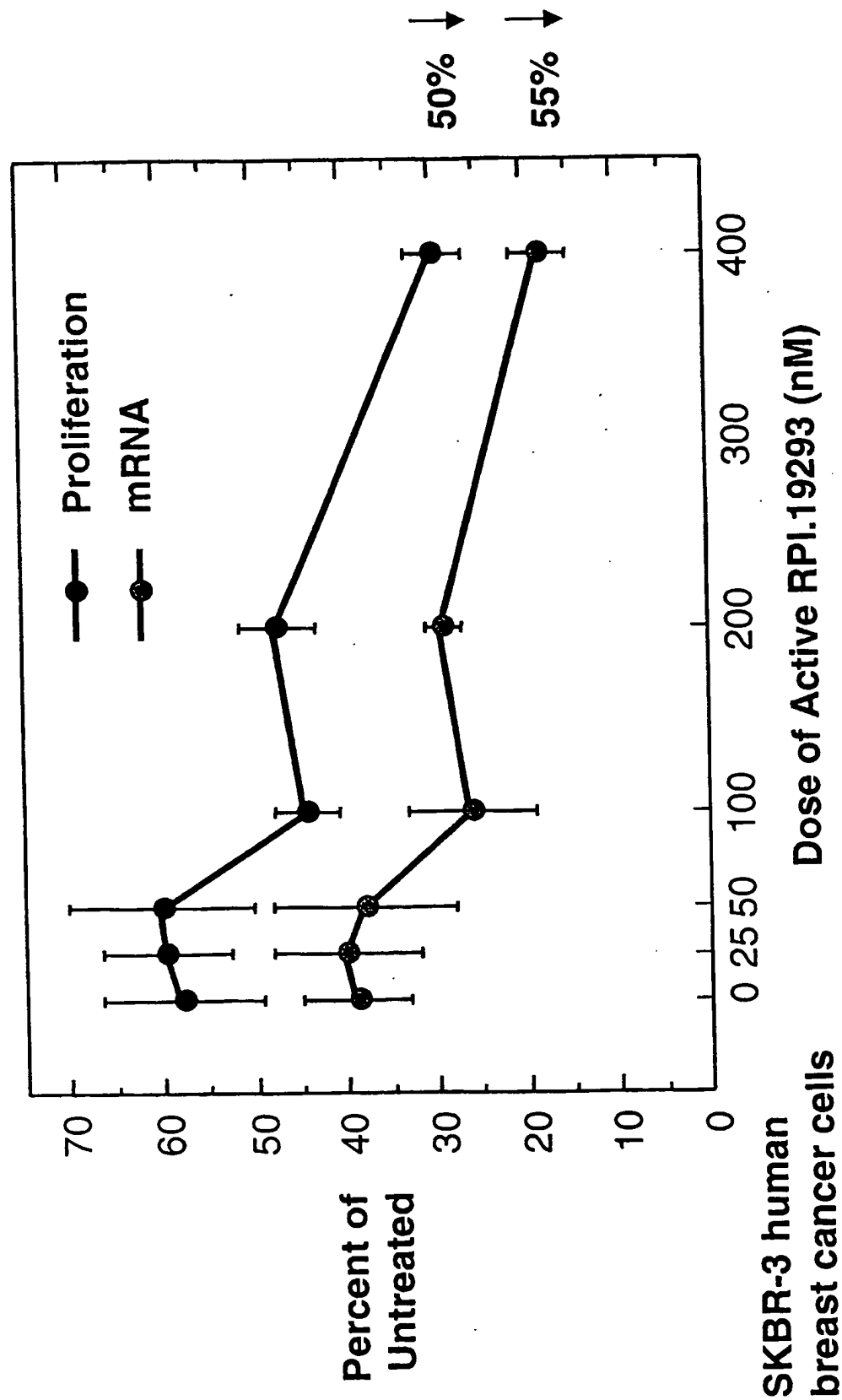
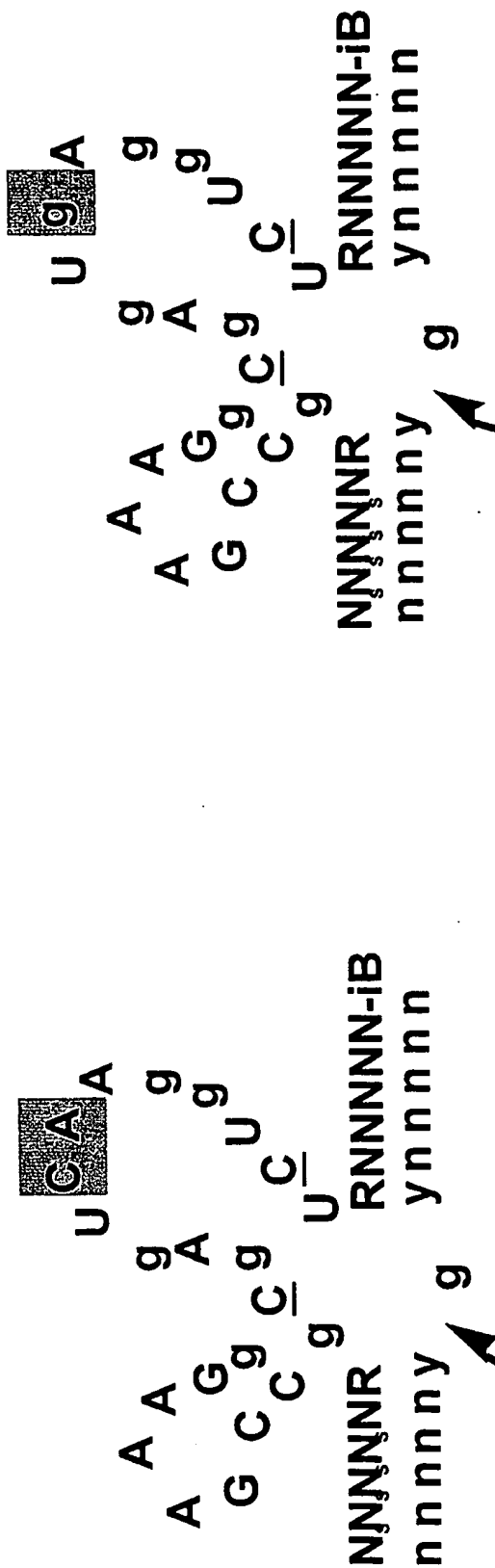


Figure 34: Zinzyme CA \rightarrow G loop (7-ribo)



Lower case = 2'-OH

Underline = 2'-NH₂

UPPER CASE = 2'-O-Me

N_s = Phosphothioate linkage

Figure 35: Screen of Zinzymes (containing ribose-G reductions) for
Anti-proliferative Activity

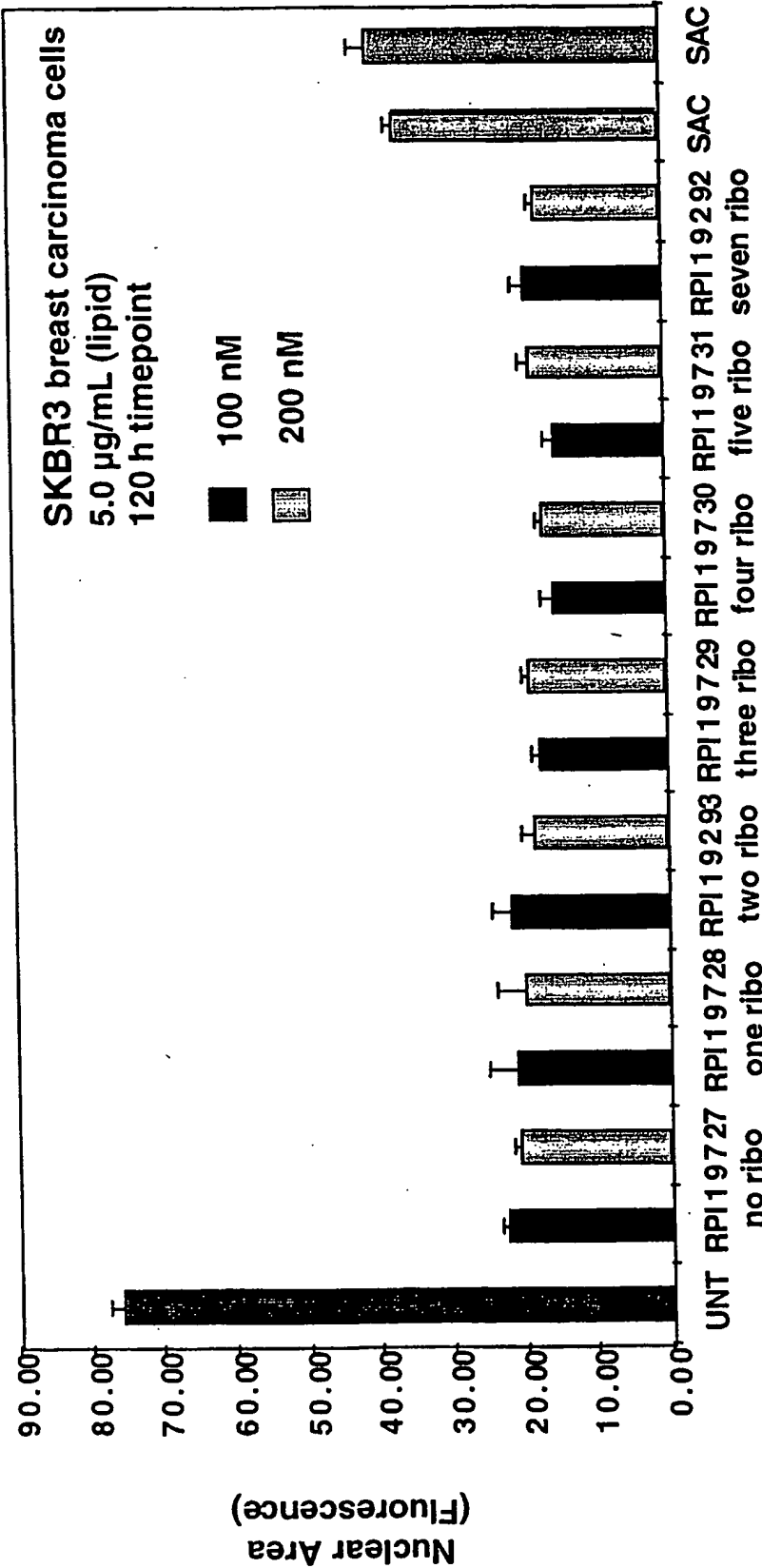
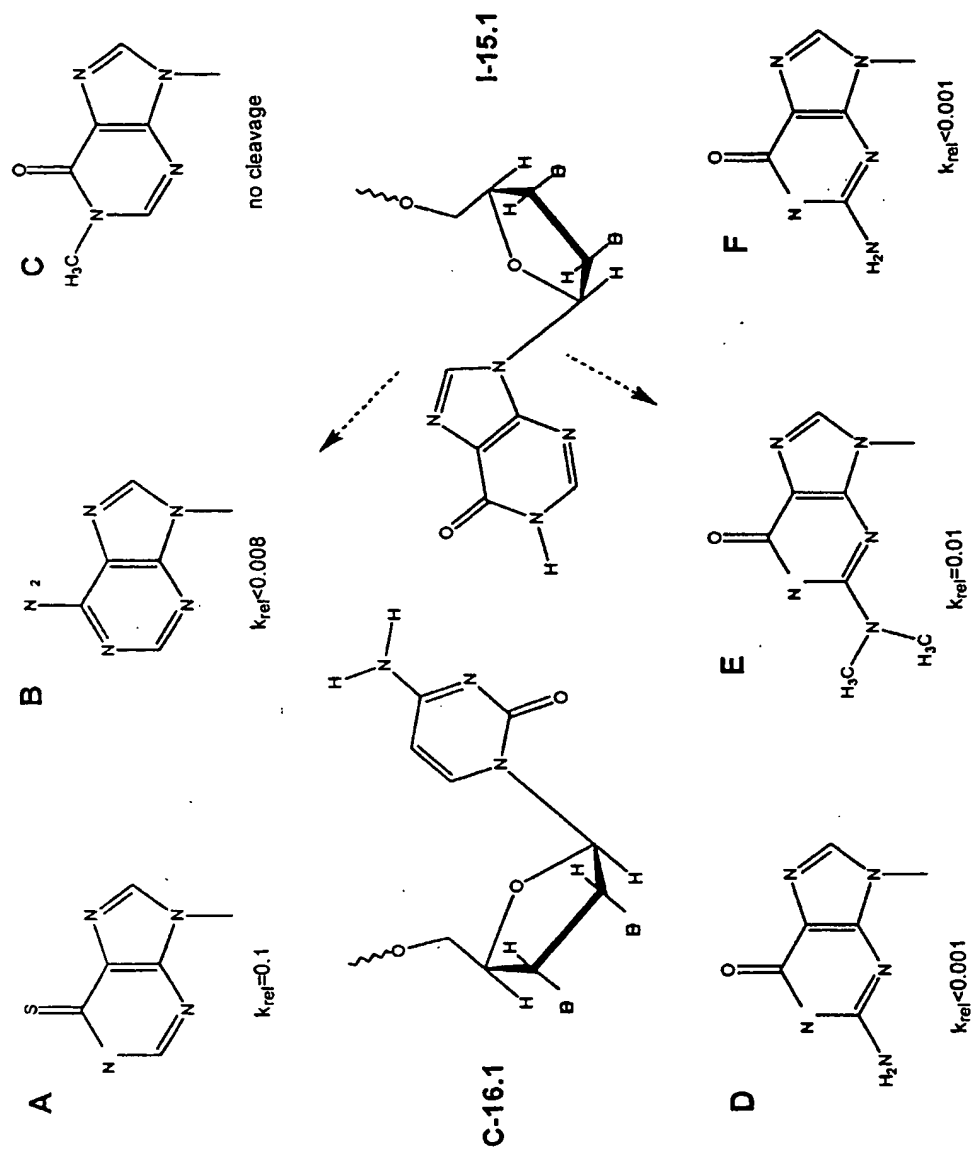
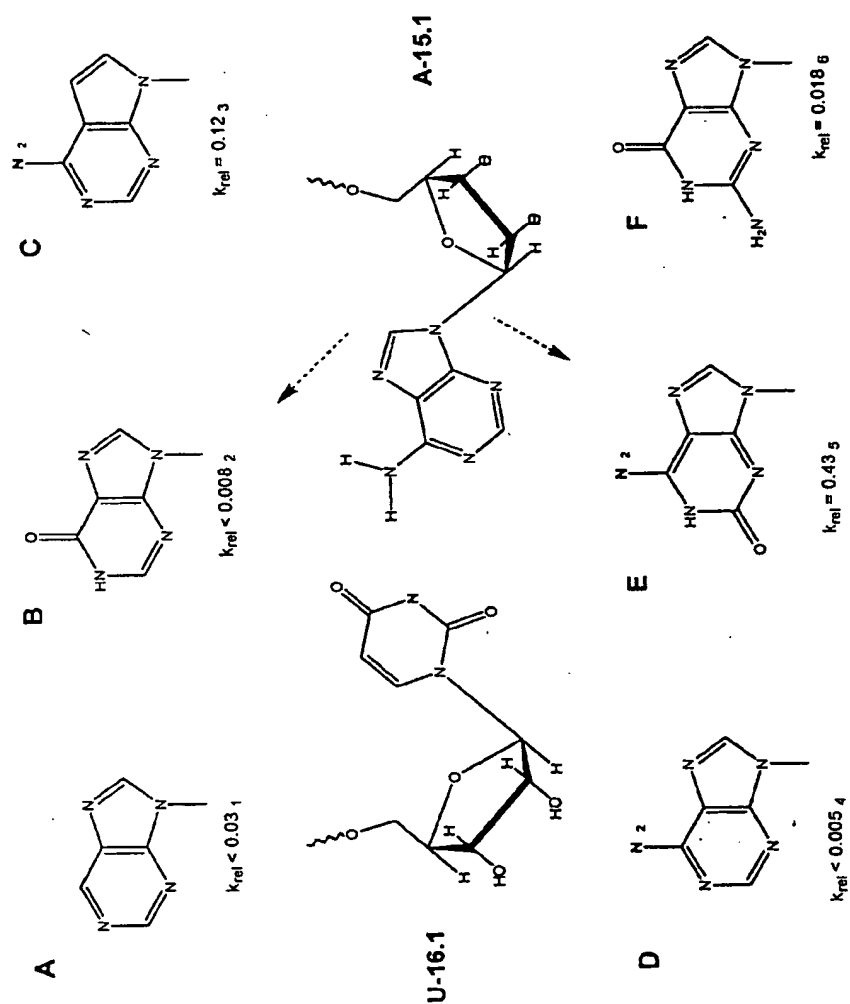


Figure 36: Effect of substitutions at NCH ribozyme position 15.1



k_{rel} values describe the cleavage rate relative to I-15.1 activity

Figure 37: Effect of substitutions at Hammerhead Ribozyme position 15.1



1. Slim and Gait, 1992, *Biochem Biophys Res Commun*, 183, 605-609.
2. Ludwig *et al.*, 1998, *Nucleic Acids Res.*, 26, 2279-2285.
3. Seela *et al.*, 1993, *Helvetica Chimica Acta*, 76, 1809-1819.
4. Seela *et al.*, 1998, *Nucleic Acids Res.*, 26, 1010-1018.
5. Ng *et al.*, 1994, *Biochemistry*, 33, 12119-26.
6. Bevers *et al.*, 1996, *Biochemistry*, 35, 6483-90.

k_{rel} values describe the cleavage rate relative to A-15.1 activity